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# Lenvatinib and immune-checkpoint inhibitors in hepatocellular carcinoma: mechanistic insights, clinical efficacy, and future perspectives

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## Abstract

Lenvatinib is a multi-target tyrosine kinase inhibitor widely used in the treatment of hepatocellular carcinoma (HCC). Its primary mechanism of action involves inhibiting signal pathways such as vascular endothelial growth factor receptors (VEGFR) and fibroblast growth factor receptors (FGFR), thereby reducing tumor cell proliferation and angiogenesis and affecting the tumor's immune microenvironment. In the treatment of liver cancer, although lenvatinib monotherapy has shown good clinical effect, the problem of drug resistance is becoming more and more serious. This resistance may be caused by a variety of factors, including genetic mutations, signaling pathway remodeling, and changes in the tumor microenvironment. In order to overcome drug resistance, the combination of lenvatinib and other therapeutic strategies has gradually become a research hotspot, and it is worth noting that the combination of lenvatinib and immune checkpoint inhibitors (ICIs) has shown a good application prospect. This combination not only enhances the anti-tumor immune response but also helps improve therapeutic efficacy. However, combination therapy also faces challenges regarding safety and tolerability. Therefore, studying the mechanisms of resistance and identifying relevant biomarkers is particularly important, as it aids in early diagnosis and personalized treatment. This article reviews the mechanisms of lenvatinib in treating liver cancer, the mechanisms and efficacy of its combination with immune checkpoint inhibitors, the causes of resistance, the exploration of biomarkers, and other novel combination therapy strategies for lenvatinib. We hope to provide insights into the use and research of lenvatinib in clinical and scientific settings, offering new strategies for the treatment of liver cancer.

**Keywords** Lenvatinib, Tyrosine kinase inhibitor (TKI), Immune-checkpoint inhibitors (ICIs), Hepatocellular carcinoma

## Introduction

### Overview of hepatocellular carcinoma (HCC)

Hepatocellular carcinoma (HCC) stands as a prevalent primary hepatic malignancy with profound global health implications. It exhibits an aggressive phenotype, high recurrence rates, and limited therapeutic avenues, culminating in a bleak prognosis characterized by a five-year overall survival rate below 20%. Chronic liver conditions such as viral hepatitis, alcohol-related liver disease, non-alcoholic fatty liver disease, and cirrhosis are closely linked to HCC, contributing to its escalating

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global incidence. Despite strides in early detection and treatment modalities, managing HCC remains a formidable challenge. Current therapeutic strategies encompass surgical resection, liver transplantation, locoregional interventions, and systemic treatments like sorafenib and lenvatinib. However, these options are encumbered by efficacy constraints, drug resistance issues, and substantial toxicities. Hence, there is an urgent call for innovative treatment paradigms to enhance outcomes for individuals afflicted with HCC.

#### **Historical context of sorafenib as a first-line therapy**

Sorafenib, a multikinase inhibitor, plays a key role in the treatment of HCC as a first-line therapy. The historical context of sorafenib as a breakthrough therapeutic option for HCC traces back to its landmark clinical trial, which demonstrated its efficacy and established it as the first systemic therapy to show a survival benefit in advanced HCC. The phase III trial, known as the SHARP trial [1], compared sorafenib to placebo in patients with unresectable HCC and demonstrated a significant improvement in median overall survival. Following the positive results of the SHARP trial, sorafenib received regulatory approval in several countries, transforming the treatment landscape for advanced HCC. Despite the subsequent emergence of other systemic therapies, sorafenib's historical significance as the first-line therapy for HCC sets the stage for subsequent developments in the field and highlights the importance of targeted therapies in improving patient outcomes. However, despite its success, sorafenib has its limitations, including modest response rates and the development of drug resistance over time.

#### **Emergence of lenvatinib as a multi-target TKI**

The development of lenvatinib as a multipotent tyrosine kinase inhibitor (TKI) is a promising alternative to these problems. Over the last few years, lenvatinib has emerged as an additional prominent multipotent tyrosine kinase (TKI) inhibitor in the first-line treatment of liver cancer (HCC). Lenvatinib's unique mechanism of action, targeting multiple kinases involved in tumor angiogenesis, proliferation, and immune regulation, has shown efficacy and safety in clinical trials. Lenvatinib acts by targeting multiple kinases involved in tumor angiogenesis, tumor proliferation, and immune regulation. Lenvatinib has been shown to be effective in REFLECT [2], a phase III trial comparing lenvatinib with sorafenib as the first-line therapy for unresectable HCC. The trial showcased non-inferiority in overall survival and demonstrated favorable outcomes in terms of progression-free survival and objective response rate, solidifying lenvatinib as a viable alternative to sorafenib. The emergence of lenvatinib as a multi-target TKI has expanded the treatment options

available to HCC patients and sparked further investigations into combination therapies, such as lenvatinib in conjunction with immune-checkpoint inhibitors, to enhance treatment responses and improve patient outcomes.

#### **Introduction to immune-checkpoint inhibitors (ICIs) in HCC**

The advent of immune-checkpoint inhibitors (ICIs) has brought significant advancements to the field of cancer immunotherapy, including their application in hepatocellular carcinoma (HCC). ICIs work by targeting key immune checkpoint molecules, such as PD-1 and PD-L1, to release the ability of the immune system to identify and eradicate cancer cells. In HCC, ICIs have shown promising clinical efficacy, and studies have demonstrated durable responses and improved survival in a subset of patients. However, the majority of HCC patients do not respond to ICIs as monotherapy, prompting the exploration of combination strategies. One such approach involves the combination of ICIs with lenvatinib, a multi-target tyrosine kinase inhibitor. Lenvatinib's anti-angiogenic properties, combined with the immunomodulatory effects of ICIs, offer a rationale for the synergistic potential of this combination therapy in HCC. Preclinical and early clinical data suggest enhanced antitumor activity and improved response rates when lenvatinib is administered in conjunction with ICIs. Understanding the mechanistic insights, evaluating the clinical efficacy, and exploring the future perspectives of the lenvatinib and ICIs combination therapy in HCC are essential for optimizing treatment outcomes and shaping the future landscape of HCC management.

This review aims to delve into the therapeutic mechanisms, clinical efficacy, and prospective directions concerning lenvatinib, particularly in combination with immune checkpoint inhibitors (ICIs), with the goal of elucidating strategies to enhance treatment efficacy and influence the evolving landscape of hepatocellular carcinoma (HCC) management.

#### **Mechanisms of action of lenvatinib in HCC**

Lenvatinib is a multi-target tyrosine kinase inhibitor (TKI) that has emerged as a promising therapeutic option in the treatment of various cancers, including hepatocellular carcinoma (HCC). It exerts its pharmacological effects by targeting multiple receptors involved in tumor angiogenesis, proliferation, and immune regulation. As a multi-target TKI, lenvatinib inhibits the activity of several key kinases involved in the development and progression of cancer. These encompass vascular endothelial growth factor receptors (VEGFR) 1–3, fibroblast growth factor receptors (FGFR) 1–4, platelet-derived growth factor receptor alpha (PDGFR $\alpha$ ), rearranged during

transfection (RET), and KIT proto-oncogene receptor tyrosine kinase (KIT). By simultaneously targeting these receptors, lenvatinib disrupts multiple signaling pathways that are critical for tumor growth and angiogenesis. The mechanisms underlying lenvatinib's anti-tumor effects involve its ability to inhibit tumor angiogenesis. By targeting VEGFRs, lenvatinib suppresses the formation of new blood vessels that supply nutrients to the tumor, thereby limiting its growth. This is especially useful for HCC, which is a highly vascularized cancer that not only receives nutrients from blood vessels but also distributes thousands of cancer cells to initiate metastasis. In addition, VEGFRs contribute to the delivery of oxygen to cancer cells. Thus, hypoxia and cell hunger are induced by lenvatinib, resulting in cell death [3]. Additionally, lenvatinib's inhibition of other receptors, such as FGFRs and PDGFR $\alpha$ , may further contribute to its anti-tumor activity by interfering with tumor cell proliferation and survival.

#### **Molecular targets and their roles in tumor proliferation and angiogenesis:**

Lenvatinib has a broad spectrum of target receptors, which allows it to disrupt multiple pathways involved in tumor proliferation and angiogenesis. This multi-pronged attack is key to its effectiveness in treating advanced hepatocellular carcinoma, where traditional treatments might fail or be insufficient.

#### ***VEGFR1-3 (Vascular endothelial growth factor receptors)***

Lenvatinib primarily inhibits tumor angiogenesis by suppressing VEGFR1-3. Tumor vascularization is intimately linked to vascular endothelial growth factor (VEGF) and its receptors (VEGFR1-3). VEGF acts directly on endothelial cells or induces angiogenesis through paracrine pathways, and it can bind to VEGFR-1 and VEGFR-2 expressed on vascular endothelial cells, triggering specific downstream signaling pathways that promote angiogenesis. Notably, VEGFR is also highly expressed in hepatocellular carcinoma [4]. Compared to normal tissue vessels, tumor blood vessels exhibit high permeability and tortuous dilation. VEGF acts on the interaction between tumor stromal cells through its specific structure, which influences the microenvironment of the tumor and plays an important role in the development of cancer.

The multiple regulatory mechanisms of VEGF give hepatocellular carcinoma a growth advantage, with VEGFR2 mediating angiogenesis and playing a central role. This makes VEGFR2 an important target for lenvatinib, which is particularly sensitive to VEGFR2-related kinases [5]. Therefore, lenvatinib can effectively inhibit tumor angiogenesis through its high affinity for VEGFR2.

VEGFR2, a type I transmembrane kinase receptor, comprises distinct domains: the extracellular domain, the transmembrane domain, and the cytoplasmic tyrosine kinase domain. Lenvatinib, akin to numerous pharmaceutical agents, exerts its effects specifically on the tyrosine kinase domain. Based on the conformation when complexed with VEGFR2, tyrosine kinase inhibitors (TKIs) are categorized into various inhibitor types (I, II, III, IV). Structural elucidation of lenvatinib demonstrates its interaction with the ATP binding site and neighboring allosteric domains in a distinct conformation (Asp-Phe-Gly, known as the "DFG-in" conformation), thereby yielding notable specificity towards kinases. Okamoto et al. suggested categorizing lenvatinib as a type V novel inhibitor [6]. In contrast, sorafenib (SOR) binds to the DFG-out state of VEGFR2, making it a type II inhibitor [7]. The differences in molecular characteristics between lenvatinib and sorafenib may account for the variations in their pharmacological activities and mechanisms of action [8]. The specific mechanisms through which lenvatinib targets VEGFR to inhibit tumor angiogenesis and affect the tumor microenvironment still require further clarification.

#### ***FGFR1-4 (Fibroblast growth factor receptors)***

Lenvatinib exhibits antitumor effects not only by inhibiting angiogenesis but also by suppressing the growth of tumor cells through the inhibition of FGFR1-4. Unlike sorafenib, lenvatinib also targets FGFR1-4, which plays a crucial role in tumor growth and serves as an important target against HCC [3]. Analysis of the co-crystal structure of the lenvatinib-FGFR1-4 complex reveals that lenvatinib also binds in a type V manner to FGFR1-4. This indicates that lenvatinib has high affinity and selectivity for FGFR in addition to VEGFR. Lenvatinib is significantly related to fibroblast growth factor (FGF) and fibroblast growth factor receptors (FGFR1-4); the FGF family promotes the proliferation and invasion of liver cancer cell lines through the FGF signaling pathway, particularly the FGF19/FGFR4 pathway [9]. Overexpression of FGF19 in liver cancer cells regulates lipid, glucose, bile acid, and energy metabolism, aiding in the survival of liver cancer cells in nutrient-limited tumor microenvironments [10]. Abnormal expression of FGF19 interacts with the  $\beta$ -Klotho co-receptor, activating the FGF19/FGFR4 signaling pathway, which promotes the proliferation, migration, and invasion of liver cancer cell lines while inhibiting apoptosis [11]. In FGF19-overexpressing HCC xenograft models, lenvatinib can inhibit the FGF signaling pathway and suppress tumor growth [8]. Clinical results also show that lenvatinib is more effective in treating HCC cases with FGF19-FGFR4 expression compared to those without this expression [12].

Research on the downstream pathways affected by lenvatinib's impact on FGFR4 indicates that lenvatinib can inhibit the phosphorylation of the downstream signaling molecule FRS2, thereby suppressing the proliferation and growth of liver cancer cells [8, 13]. Additionally, lenvatinib can reduce xCT expression by inhibiting FGFR4, ultimately inducing lipid ROS accumulation and leading to ferroptosis in liver cancer cells [14]. In the context of the immune microenvironment, empirical investigations conducted both in vitro and in vivo have revealed that lenvatinib diminishes tumor PD-L1 expression and inhibits Treg cell differentiation through FGFR4 blockade, consequently augmenting the effectiveness of anti-PD-1 immunotherapy. These findings furnish critical theoretical underpinnings for the synergistic application of lenvatinib in conjunction with anti-PD-1 therapy [15].

In addition to FGFR4, a preclinical study using an HCC PDX model indicated that models with high FGFR1 expression respond better to lenvatinib than to sorafenib. Prolonged exposure to sorafenib can lead to upregulation of FGFR1, causing HCC cells to develop resistance to sorafenib. Therefore, lenvatinib may potentially overcome this sorafenib resistance by blocking FGFR1 [8]. Lenvatinib can also reduce cancer stem-like cells (CSCs) in HCC by inhibiting FGFR1-3 signaling pathways.

As two important target receptors of lenvatinib, VEGFR and FGFR are closely interconnected after lenvatinib treatment. Suppression of FGFR activity triggers a compensatory activation of the EGFR–PAK2–ERK5 signaling cascade, which can be effectively attenuated by targeting EGFR, thereby interrupting this signaling pathway. Therefore, combined inhibition of FGFR and EGFR makes liver cancer more sensitive to lenvatinib [16]. Additionally, the FGF-FGFR-MAPK axis mediates the survival of HCC cells under nutrient-deprived conditions due to vascular suppression. Consequently, using drugs like lenvatinib to dual-target both VEGFR and FGFR provides enhanced antitumor activity against HCC with activated FGF signaling pathways compared to therapies that only target VEGF signaling [17]. These findings suggest that lenvatinib significantly enhances the antitumor response in liver cancer compared to sorafenib, due to its simultaneous action on both VEGFR and FGFR.

**Other target receptor: PDGFR $\alpha$  (Platelet-derived growth factor receptor), RET (Rearranged during transfection), and KIT (KIT proto-oncogene receptor tyrosine kinase)**

Compared to the extensively studied target receptors VEGFR and FGFR, research on the remaining receptors, PDGFR $\alpha$ , RET, and KIT, is relatively limited. PDGFR is a receptor tyrosine kinase widely distributed on the surface of various cells, existing in two structurally similar subtypes: PDGFR $\alpha$  and PDGFR $\beta$ . Dysregulated activation of

the PDGF/PDGFR pathway has the potential to stimulate angiogenesis, facilitating the development of neovascularization within tumors, while concurrently influencing tumor cell proliferation and migration through direct or indirect mechanisms. The RET protein, encoded by the RET gene, belongs to the receptor tyrosine kinase family. Pathogenic mutations—either mutations or rearrangements—of the RET gene can lead to its abnormal activation, resulting in tumor cell proliferation, differentiation, and migration. However, research on lenvatinib targeting RET has largely been limited to thyroid cancer, likely due to the high mutation rate of RET in that context, drawing researchers' attention [18]. Both in vitro and in vivo studies have shown that lenvatinib can inhibit RET phosphorylation and subsequently suppress RET-ERK signaling in cases with genetic alterations in RET (such as activating mutations or gene rearrangements) [19, 20], ultimately affecting the antitumor process. Kato et al. reported that among 4871 cases of other cancer types analyzed for RET mutations and amplifications, only 88 cases (1.8%) were observed, with mutations found in just one of 44 HCC cases [21]. Therefore, the frequency of RET mutations in HCC is low, and the inhibitory effect of lenvatinib on RET may be limited to certain cases. A comprehensive understanding of the mechanisms by which lenvatinib affects the PDGFR, RET, and KIT pathways is crucial for revealing its therapeutic mechanisms. Future research is anticipated to delve deeper into the specific roles of these pathways and provide stronger support for personalized treatment strategies.

**Impact on the tumor immune microenvironment:**

Lenvatinib not only inhibits tumor cell growth in the treatment of hepatocellular carcinoma (HCC) but also significantly impacts the tumor immune microenvironment. It alters the immune status of the tumor microenvironment by regulating the activity and distribution of immune cells, thereby enhancing the body's immune response against the tumor. This immunomodulatory effect may contribute to improved treatment outcomes and better prognosis for HCC patients. Additionally, understanding lenvatinib's influence on the immune microenvironment will provide new insights for future combination immunotherapy strategies.

Lenvatinib regulates the tumor immune microenvironment in hepatocellular carcinoma (HCC) through multiple mechanisms, promoting immune cell infiltration and function, thus enhancing treatment efficacy. It plays a significant role in the HCC immune microenvironment by targeting the VEGFR pathway. Research shows that lenvatinib downregulates the VEGFR pathway and promotes the infiltration of GZMK+CD8 T cells in the tumor microenvironment, mediated by macrophage-derived

CXCL9. However, the recruited T cells may lose their original antitumor activity due to the physical barriers of the extracellular matrix (ECM). Therefore, compounds targeting the ECM could facilitate the penetration of cytotoxic GZMK+CD8 T cells into HCC cell areas, ultimately improving drug efficacy [22]. Additionally, lenvatinib can impact the PI3K/Akt/mTOR signaling pathway in PD1+CD8+T cells by targeting VEGFR2 on these cells, influencing their differentiation, promoting the self-proliferation of TCF1+PD1+Ki-67+TIM3-T progenitor cells, and enhancing the tumor-killing function of intermediate TCF1-PD1+Ki-67-TIM3-T cells [23]. Beyond targeting VEGFR, lenvatinib also participates in the HCC tumor immune process by targeting FGFR [8]. Moreover, lenvatinib affects metabolic pathways that influence the immune microenvironment. Research by Sun et al. indicates that lenvatinib targets TET2 synthesis to increase the levels of nicotinamide adenine dinucleotide (NAD<sup>+</sup>), enhances the secretion of inosine, and inhibits the secretion of nicotinamide, 5-hydroxy-L-tryptophan, and quinolinic acid, thereby promoting macrophage proliferation and migration and facilitating M1 polarization [24]. Under the influence of lenvatinib, the properties of immune cells also change. Besides inducing macrophage polarization, lenvatinib significantly reduces the expression of PD-1 and TIM-3 on CTLs, as well as TIM-3 and CTLA-4 on Treg cells [25]. This provides a theoretical basis for the combined application of lenvatinib with immune checkpoint inhibitors, potentially further enhancing antitumor immune responses and improving treatment outcomes.

There is extensive research on the components of immune cells and immune factors in the tumor microenvironment after lenvatinib treatment. Following treatment, the frequency of T helper (Th) cells and regulatory T (Treg) cells decreases, while cytotoxic T lymphocytes (CTL) significantly increase. The CTL/Treg ratio has been identified as a new prognostic indicator for HCC patients [25]. Additionally, the ratios of monocytes and macrophages, as well as neutrophils and lymphocytes, are significantly reduced, while the numbers of CD4+T cells, CD8+T cells, and Th1 cells increase [26–28]. The cytokine profile shows an increase in IL-2, IL-5, and IFN- $\gamma$ , while IL-6, IL-10, TNF- $\alpha$ , and TNF- $\beta$  levels decrease [25]. These changes reflect a re-regulation of the immune system and are closely associated with patient prognosis. Future research will further explore the roles of these immune cells and factors, providing more evidence for personalized treatment strategies.

Overall, lenvatinib exhibits dual efficacy by suppressing tumor cell proliferation in hepatocellular carcinoma (HCC) and substantially altering the tumor immune microenvironment, thereby augmenting the

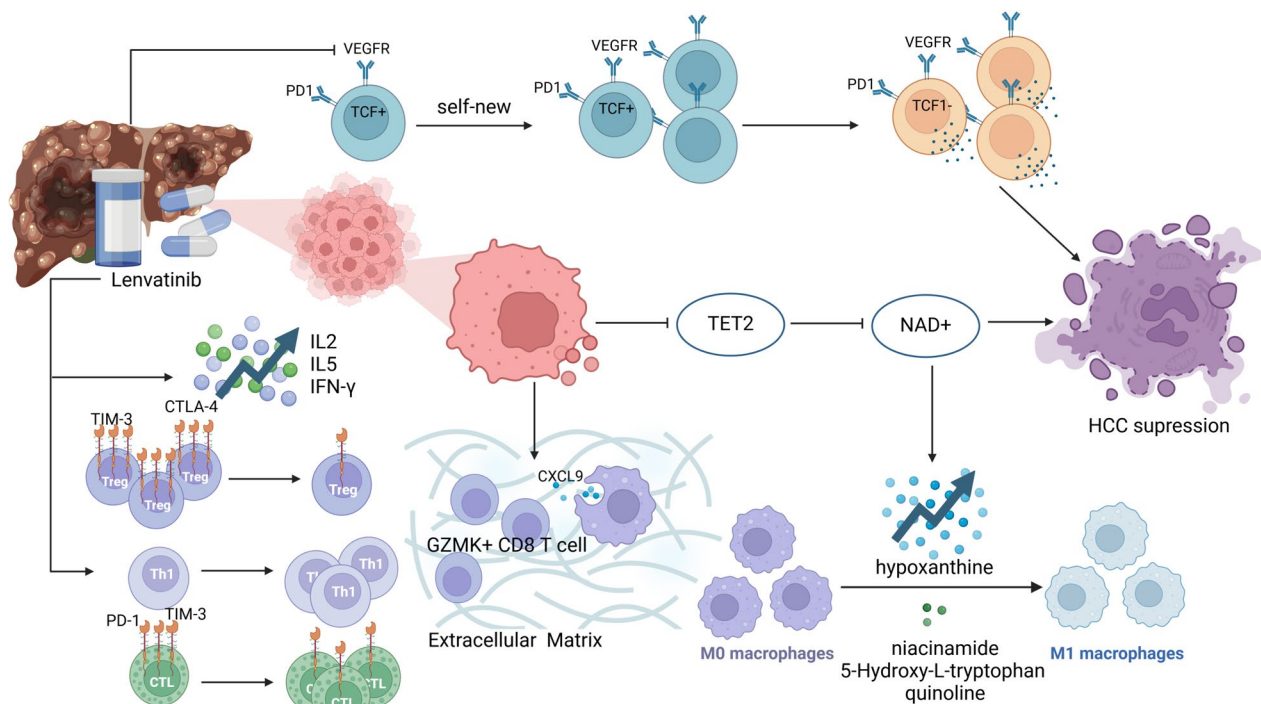
host immune response. This immunomodulatory effect may improve treatment efficacy and patient prognosis. Further research on lenvatinib's impact on the immune microenvironment will provide an important theoretical foundation for future combination immunotherapy strategies (Fig. 1).

#### **Antitumor proliferation and immunomodulatory activity in preclinical models**

Researchers have investigated the cellular-level mechanism of action of lenvatinib, validating its antitumor proliferation and immunomodulatory activity through cell experiments. Subsequently, the research progressed to preclinical models, assessing the drug's efficacy and safety in more complex biological systems. This process not only enhances the credibility of the research but also lays a solid foundation for future clinical applications, allowing the findings to better inform the development of actual treatment plans. Encouragingly, preclinical models also demonstrate that lenvatinib has significant antitumor proliferation and immunomodulatory activity in the treatment of liver cancer.

Lenvatinib has demonstrated impressive performance in preclinical models, particularly in anti-angiogenesis. The strong activity of lenvatinib against VEGF signaling in xenograft models of HCC with overexpressed VEGF provides the basis for its antitumor and antiangiogenic effects in vascularized HCC models [29]. Additionally, evidence shows that lenvatinib's ability to inhibit tumor growth in mice is dose-dependent, meaning that as the dose of lenvatinib increases, tumor vascularity and microvascular density significantly decrease, while necrosis rates markedly increase [30]. Interestingly, some studies have approached this from an imaging perspective. Results indicate that early in lenvatinib treatment, CT scans show no significant decrease in tumor blood vessel volume density, with a notable decline in TBV (tumor blood volume) density occurring by day fourteen. This suggests that the normalization of TBV, which begins early in treatment, is a unique feature of lenvatinib therapy. It appears that lenvatinib does not affect existing vascular structures but rather induces a reduction in vessel diameter in terms of TBV normalization [31]. This research helps to better understand how lenvatinib influences vascular changes.

In addition to affecting tumor angiogenesis, preclinical models treated with lenvatinib have also shown changes in the immune microenvironment. In an orthotopic rat model carrying liver cancer cells, administering 5 mg/kg of lenvatinib demonstrated that lenvatinib had no direct effect on macrophages, but instead led to their polarization towards the M1 phenotype. This polarization



**Fig. 1** Effect of lenvatinib on the tumour immune microenvironment

improves the tumor immune microenvironment and contributes to antitumor effects [32].

Many preclinical studies have compared lenvatinib and sorafenib. In HCC patient-derived xenograft (PDX) models, research has confirmed that lenvatinib exhibits superior anti-angiogenic capabilities compared to sorafenib. FGF, a pro-angiogenic factor, can induce escape from anti-VEGF therapy, while lenvatinib targets FGFR, a capacity that sorafenib lacks [8]. In the context of FGF-induced angiogenesis, examination through histological analysis of xenograft tumors overexpressing FGF19 in mice treated with lenvatinib demonstrated that FGF facilitates the survival of hepatocellular carcinoma (HCC) cells characterized by activated FGF signaling pathways within tumor microenvironments deprived of adequate nutrients, operating through the MAPK cascade [17]. Moreover, in murine models harboring subcutaneously implanted hepatocellular carcinoma (HCC), the administration of 3 mg/kg of lenvatinib demonstrated a significant reduction in microvascular density and the normalization of tumor vasculature compared to treatment with 50 mg/kg of sorafenib. These findings indicate that lenvatinib expedites tumor vascular normalization and improves the intra-tumoral microenvironment more rapidly and effectively than sorafenib [33].

These studies indicate that lenvatinib not only suppresses tumor cell growth but also effectively regulates

immune cells in the tumor microenvironment, boosting the host's immune response. By thoroughly analyzing the results from these models, researchers can attain a more comprehensive comprehension of lenvatinib's mechanisms of action, providing crucial evidence for its efficacy in clinical applications.

### Immune-checkpoint inhibitors in HCC

Immune checkpoint inhibitors in oncology are a class of innovative drugs used for cancer treatment. They activate the patient's immune system to attack and suppress tumor growth by blocking the inhibitory signals between tumor cells and the immune system. These inhibitors primarily target immune checkpoints, which are key molecules that regulate immune responses. Immune checkpoint inhibitors, such as PD-1 and PD-L1 inhibitors, have demonstrated significant efficacy across various cancer types. For example, in melanoma, non-small cell lung cancer, and renal cell carcinoma, immune checkpoint inhibitors have become part of standard treatment, significantly improving patient survival rates and quality of life. Additionally, ICIs have broadened treatment options, maintained treatment durability, and provided new approaches for personalized therapy.

Typical immune checkpoint inhibitors (ICIs) encompass anti-programmed cell death-1 (PD-1) and programmed cell death ligand 1 (PD-L1) therapeutics,

exemplified by nivolumab, pembrolizumab, durvalumab, and atezolizumab, along with anti-cytotoxic T lymphocyte-associated protein 4 (CTLA-4) agents such as tremelimumab and ipilimumab [34]. Lymphocyte activation gene 3 (LAG-3) is the most mature immune checkpoint studied after CTLA-4 and PD-1/PD-L1, with LAG-3 antibodies like Opdualag and Relatlimab already approved for clinical anti-tumor therapy. Additionally, T cell immunoglobulin and mucin domain-containing protein 3 (TIM-3), CD47, T cell immunoglobulin and ITIM domain protein (TIGIT), and V-domain Ig suppressor of T-cell activation (VISTA) are also under extensive research and clinical trials, which is significant for overcoming acquired resistance to immune checkpoint inhibitors and preventing the emergence of intrinsic resistance mechanisms.

### Mechanisms of action of ICIs

#### PD-1/PD-L1

Programmed cell death protein 1 (PD-1) is a negative co-stimulatory receptor expressed on activated T cells, natural killer (NK) cells, and monocytes. Its ligand, programmed death-ligand 1 (PD-L1), engages PD-1 on T cells, triggering tyrosine phosphorylation within the immune receptor tyrosine motif of PD-1. This event recruits Src homology region 2-containing protein tyrosine phosphatase 2 (SHP-2), leading to the dephosphorylation of downstream protein kinases like spleen tyrosine kinase (Syk) and phosphatidylinositol 3-kinase (PI3K). Consequently, this process disrupts signaling pathways involving protein kinase B (PKB or AKT) and extracellular regulated protein kinase (ERK), impeding gene transcription and cytokine production necessary for T cell activation, ultimately imposing negative regulation on T cell function [35]. Tumor cells exhibiting elevated PD-L1 levels can interact with PD-1 on tumor-specific T cells, impeding T cell activation, proliferation, and cytokine release, thereby facilitating immune evasion by the tumor. Consequently, interruption of the PD-1/PD-L1 signaling axis can revive T cell cytotoxicity and instigate inherent anti-tumor immune responses, thereby manifesting anti-tumor properties.

Since 2012, following the favorable safety profile and anti-tumor efficacy demonstrated by PD-1 monoclonal antibodies in phase I/II clinical trials, multiple phase III clinical trials have been initiated targeting various malignancies. Presently, a total of 10 PD-1 monoclonal antibodies and 3 PD-L1 monoclonal antibodies have received approval for treating 11 cancer types [35], including liver cancer.

Although PD-1/PD-L1 antibodies exhibit significant anti-tumor activity across various malignancies, their response rates are relatively low in most tumors.

Developing combination strategies with PD-1/PD-L1 antibodies is one approach to enhance the response rates of PD-1/PD-L1 monotherapy. Clinical trials combining PD-1/PD-L1 therapies with other anti-tumor agents are underway, including other immunotherapies (such as other ICIs, oncolytic viruses, stimulator of interferon genes (STING) agonists, and chimeric antigen receptor T-cell (CAR-T) therapy), radiotherapy, chemotherapy, and molecularly targeted therapies. However, currently, FDA-approved PD-1/PD-L1 combination strategies are limited to combinations with chemotherapy, vascular endothelial growth factor (VEGF) targeted agents, and CTLA-4 monoclonal antibodies.

#### CTLA-4

CTLA-4 is widely expressed in activated T cells. The binding of the T cell surface receptor CD28 to CD80/CD86 on antigen-presenting cells (APCs) generates a co-stimulatory signal, which, along with the signal from the antigen-specific T cell receptor (TCR) recognizing major histocompatibility complex (MHC), constitutes the dual signal necessary for T cell activation. CTLA-4 shares a high degree of homology with the extracellular domain of CD28, and compared to CD28, CTLA-4 has a higher affinity for CD80/CD86. This high affinity allows CTLA-4 to antagonize the interaction between CD28 and its ligands, thereby reducing the generation of co-stimulatory signals and inhibiting T-cell activation. Additionally, CTLA-4 is also expressed on regulatory T cells (Tregs), where it promotes the internalization of CD80/CD86 on APCs through trans-endocytosis, reducing their expression levels and further enhancing immunosuppressive functions. Therefore, targeting and blocking CTLA-4 can relieve T cell immune suppression and allow for T cell activation, thereby facilitating tumor cytotoxicity.

Recent investigations indicate that antibody-dependent cell-mediated cytotoxicity (ADCC) may represent a fundamental mechanism through which ipilimumab mediates its anti-neoplastic actions. Research indicates that ADCC, mediated by the Fc region, can target and eliminate Tregs that express high levels of CTLA-4, thereby relieving the immune suppression caused by these Tregs. Consequently, enhancing Fc-mediated ADCC has become an important direction in the development of CTLA-4 monoclonal antibodies.

One phase III clinical trial of tremelimumab [36] (Clinical Trial Number: NCT03298451) showed significant improvement in overall survival (OS) when combined with durvalumab in patients with liver cancer. Based on this study, tremelimumab has received FDA priority review and is expected to become the second approved CTLA-4 monoclonal antibody. Besides these drugs, numerous CTLA-4 antibody candidates have entered

clinical and preclinical research stages, focusing primarily on the efficacy and safety of the monoclonal antibodies.

In terms of combination therapy, CTLA-4 monoclonal antibodies target the early stages of anti-tumor immunity and have a long-lasting effect. When combined with PD-1/PD-L1 monoclonal antibodies, which act during the T cell effector phase, this approach can significantly improve response rates and enhance anti-tumor efficacy. The amalgamation of the anti-PD-1 antibody nivolumab with the anti-CTLA-4 antibody ipilimumab was initially evaluated in the phase I/II clinical trial CheckMate-040 [37] (NCT01658878) in liver cancer. Compared to monotherapy, the nivolumab and ipilimumab combination appears to be safe and effective.

### **LAG-3**

LAG-3 is another important immune checkpoint following CTLA-4 and PD-1/PD-L1. It is expressed on the surface of various lymphocytes and is structurally similar to CD4, with a stronger affinity for MHC class II. LAG-3 can stably bind to peptide-MHC class II complexes, inhibiting T-cell activation. Additionally, fibrinogen-like protein 1 (FGL1) has been identified as a ligand for LAG-3. Blocking the FGL1-LAG-3 interaction can also enhance tumor immunity and inhibit tumor growth in tumor-bearing mice.

Relatlimab is the first LAG-3 monoclonal antibody developed for clinical use. Although LAG-3 antibody monotherapy has good safety, its diverse ligands and mechanisms of action limit its anti-tumor efficacy. Current clinical studies indicate that the combination of LAG-3 with PD-1/PD-L1 antibodies holds significant promise in terms of efficacy and safety. Several other LAG-3 antibodies, such as GSK2831781, HLX26, and Sym022, are currently in clinical trials.

### **Other ICIs**

TIM-3 is also a negative co-stimulatory molecule, widely expressed on various immune cells. Currently, four ligands have been identified for TIM-3, including galectin-9 (Gal-9), carcinoembryonic antigen-related cell adhesion molecule 1, high mobility group protein B1, and phosphatidylserine, with Gal-9 being the primary ligand. TIM-3 mediates dysfunction and exhaustion of TIM-3+ T cells through its interaction with Gal-9. Currently, antibody drugs targeting TIM-3, such as Sym023, INCAGN2390, and LY3321367, are undergoing clinical trials as monotherapies for advanced solid tumors and lymphomas (NCT03489343, NCT03652077, NCT04443751). LY3321367 has shown good safety in phase I trials, but its anti-tumor activity is limited. Additionally, studies indicate that the development of resistance to PD-1 antibodies is associated

with the upregulation of TIM-3. A clinical trial of TIM-3 and PD-1 antibody combination therapy for advanced solid tumors showed that the combination group not only had good tolerability but also demonstrated improved anti-tumor activity. Research on the combination of TIM-3 and PD-1 antibodies is currently being widely conducted.

CD47 is widely expressed in normal cells and is overexpressed in various types of solid tumors and hematological malignancies. The regulation of tumor immunity by CD47 primarily occurs through its interaction with signal regulatory protein  $\alpha$  (SIRP $\alpha$ ) on macrophages, inhibiting the clearance of tumor cells by these immune cells. To enhance the efficacy of CD47 antibodies, numerous clinical trials combining CD47 antibodies with other immune checkpoint inhibitors (ICIs) are currently being conducted, pending further research for confirmation.

TIGIT is primarily expressed in T cells and NK cells, with three ligands: CD155, CD112, and CD113 [38]. Among these, TIGIT has a higher binding affinity for CD155 compared to CD112 and CD113. The interaction between TIGIT and CD155 inhibits the activity of T cells and NK cells, leading to reduced antigen presentation by dendritic cells and decreased secretion of anti-inflammatory cytokines, which ultimately impairs T cell activation. Research has shown that TIGIT antibody monotherapy is insufficient to suppress tumor growth in mice, but administering TIGIT monoclonal antibodies before tumor formation can delay tumor development and metastasis. Additionally, studies indicate that combining TIGIT antibodies with PD-1/PD-L1 antibodies can enhance the anti-tumor effects of TIGIT antibodies. Currently, several TIGIT inhibitors are in clinical studies but none have been approved for marketing. Among them, only four TIGIT antibodies have reached phase III clinical trials: tiragolumab, vibostolimab, ociperlimab, and domvanalimab.

VISTA exhibits sequence similarity with PD-L1 and PD-L2 and demonstrates notable expression in myeloid-derived suppressor cells and various immune cell populations. Currently, two immunosuppressive ligands have been confirmed for VISTA: V-set and immunoglobulin domain containing 3 (VSIG3) and P-selectin glycoprotein ligand 1 (PSGL-1). At physiological pH, VISTA interacts with VSIG3, while at acidic pH, VISTA-expressing cells can bind to PSGL-1 on T cells. Both interactions suppress T cell function. Although antibodies targeting VISTA have shown significant therapeutic effects in various preclinical tumor models, there are currently no approved antibody drugs on the market. Most effective VISTA treatment models also rely on combination approaches.



### Specific ICIs used in HCC

In the era of immunotherapy, the use of immune checkpoint inhibitors (ICIs) in liver cancer is of significant importance and drives the development of personalized and precision medicine. ICIs have also been tested in HCC patients, with nivolumab and pembrolizumab (both anti-PD-1 antibodies) approved for second-line treatment.

In 2012, a phase I/II clinical trial, CheckMate 040, was initiated to evaluate the efficacy and safety of nivolumab, either as a monotherapy or in combination, for second-line treatment of advanced hepatocellular carcinoma (HCC) [39]. Following the release of preliminary results from CheckMate 040 in 2017, the FDA approved nivolumab for second-line treatment of HCC, making it the first immunotherapy approved for this indication. A study comparing nivolumab monotherapy to sorafenib monotherapy for first-line treatment of advanced HCC showed that nivolumab did not significantly improve overall survival compared to sorafenib, but clinical activity and good safety were observed in advanced HCC patients [40]. Therefore, nivolumab may be considered a treatment option for patients contraindicated or at significant risk with tyrosine kinase inhibitors and anti-angiogenic agents. Regarding the long-term effects of nivolumab monotherapy, a five-year follow-up demonstrated that it provided durable clinical benefits and manageable safety for both treatment-naïve and sorafenib-treated advanced HCC patients, with the five-year survival rate for advanced HCC increasing from 3 to 12%–14% [41].

KEYNOTE-224 was a pioneering study of pembrolizumab monotherapy for second-line treatment of advanced HCC [42], and due to its outstanding results, pembrolizumab was approved by the FDA. Second-line treatment is the primary focus for pembrolizumab. Following the KEYNOTE-224 study, researchers initiated two large multicenter phase III trials: KEYNOTE-240 and KEYNOTE-394. In the global KEYNOTE-240 study, the primary endpoint did not achieve statistical significance [43]. However, the KEYNOTE-394 trial in the Asian population met its primary endpoint [44], making it the first phase III clinical trial of an immune monotherapy to achieve positive results in liver cancer. The trial demonstrated positive outcomes for overall survival (OS), progression-free survival (PFS), and overall response rate (ORR). The OS in the pembrolizumab group reached 14.6 months, representing the longest OS data for immune monotherapy in liver cancer to date, while the duration of response (DOR) was 23.9 months, offering hope for prolonged survival in liver cancer patients.

In numerous instances, sole targeting of immune checkpoints has demonstrated inadequacy, given that

these checkpoints represent only a fraction of the complexities inherent in the immune system and tumor microenvironment. As the application of ICIs expands, resistance phenomena have gradually emerged, with many patients experiencing tumor recurrence or progression after initial treatment. This resistance mechanism may be associated with alterations in the tumor microenvironment, activation of immune evasion pathways, and the genomic diversity of tumor cells. As a result, a comprehensive investigation into resistance mechanisms and the establishment of strategies to overcome them are crucial for enhancing the therapeutic efficacy of immune checkpoint inhibitors. The realm of combination therapies based on immune checkpoint inhibitors is progressively gaining traction, supplanting tyrosine kinase inhibitor monotherapy in advanced hepatocellular carcinoma (HCC). Nevertheless, challenges such as resistance, the identification of predictive biomarkers for treatment response, and potential adverse effects in combined therapies persist. The development of safer and more effective combination treatment protocols for advanced HCC is paramount, underscoring the need for further research endeavors.

### Synergy between lenvatinib and ICIs:

Not all HCC patients, especially those awaiting liver transplantation, respond to ICI immunotherapy. More importantly, monotherapy with immunotherapy does not significantly improve overall survival (OS). Given this data, researchers are exploring more effective combination therapies for HCC, including the combined use of ICIs and TKIs [34].

### Role of lenvatinib in modulating the tumor microenvironment to enhance ICI efficacy

The mechanism by which lenvatinib enhances ICI efficacy is closely related to its target pathways. Lenvatinib's modulation of the VEGF pathway can diminish the activity of cytotoxic T cells and dendritic cells, while facilitating the infiltration of immunosuppressive cell populations such as regulatory T cells (Tregs), myeloid-derived suppressor cells (MDSCs), and M2 macrophages, thereby exerting a direct immunosuppressive impact within the tumor microenvironment [45]. Additionally, lenvatinib can induce the formation of the NRP-1-PDGFR $\beta$  complex and activate the Crkl-C3G-Rap1 signaling pathway in endothelial cells, further optimizing the immature and dysfunctional tumor vasculature. This enhances the infiltration of immune cells, primarily CD8+ T cells, reshaping the tumor microenvironment (TME) and increasing sensitivity to immune checkpoint blockade (ICB) therapies [46]. Moreover, the inhibition of VEGFR-2 in endothelial cells triggers the expression

of PD-L1 in hepatocellular carcinoma (HCC) cells in a paracrine fashion, partially mediated by interferon- $\gamma$  expression. Concurrently, VEGFR-2 blockade enhances PD-1 expression in CD4+ cells infiltrating the tumor. In the presence of anti-PD-1 therapy, CD4+ cells facilitate the normalization of vascular structures when exposed to anti-VEGFR-2 antibodies during anti-angiogenic treatment, laying the groundwork for potential combined therapeutic approaches [47].

Similarly, the FGFR-FGF pathway is actively involved in the mechanism by which lenvatinib enhances ICI efficacy. In vitro and in vivo experiments have demonstrated that lenvatinib improves anti-PD-1 efficacy by reducing tumor PD-L1 levels and Treg differentiation through FGFR4 blockade. The expression levels of FGFR4 and Treg infiltration in tumors can serve as biomarkers for screening HCC patients for combination therapy with lenvatinib and anti-PD-1 [15]. Activation of FGFR signaling inhibits the JAK/STAT signaling pathway stimulated by IFN- $\gamma$ , reducing the expression of its target genes, including B2M, CXCL10, and PD-L1 [48]. Inhibiting FGFR with lenvatinib can enhance anti-tumor immunity and the activity of anti-PD-1 antibodies. Additionally, lenvatinib's suppression of FGFR signaling can increase the number of PD-L1-positive cells by downregulating lipid metabolism-related genes, transforming the tumor immune microenvironment (TIME) into a hot immune state [49].

Lenvatinib also has potential effects on the inhibition of other targets such as rearranged during transfection (RET) and platelet-derived growth factor (PDGF), carrying additional immunological and molecular significance [45]. Furthermore, lenvatinib can restore sensitivity to ICIs by blocking the PKC $\alpha$ /ZFP64/CSF1 axis, thereby reshaping the tumor microenvironment [50].

By enhancing the efficacy of ICIs within the tumor microenvironment of HCC, the combination of lenvatinib and immune checkpoint inhibitors has become an important treatment strategy. This combination can improve the tumor microenvironment, promote the infiltration and activity of immune cells, and enhance the body's immune response to tumors. This combined treatment approach offers new hope for HCC patients, particularly those who do not respond well to traditional therapies, further advancing the development of personalized medicine.

#### ***Preclinical data showing enhanced anti-tumor immune response***

Lenvatinib and ICI combination therapy have shown significant superiority in preclinical liver cancer animal models. In murine models, the combined treatment exhibited tumor regression and quicker response

durations in contrast to monotherapy ( $P < 0.001$ ). Monotherapy involving anti-PD-1 triggered infiltration of dendritic cells and T cells, whereas lenvatinib decreased the percentage of regulatory T cells (Tregs). Nevertheless, solely the combined therapy notably suppressed immunosuppressive signaling associated with the TGF $\beta$  pathway and fostered an immune-active microenvironment ( $P < 0.05$ ) [45]. In a study conducted by Chen et al., it was observed that the co-administration of lenvatinib with siRNA targeting PD-L1 effectively impeded tumor progression in H22 tumor-bearing mice by impeding tumor cell proliferation and fostering apoptotic processes. This combined therapeutic regimen synergistically downregulated the expression of vascular endothelial growth factor (VEGF) and PD-L1, facilitating heightened infiltration of T cells within tumor tissues and an increased T cell population in the spleen. Furthermore, the combined treatment amplified the presence of granzyme B-positive T cells, suggesting a notable reinforcement of the mice's anti-tumor immune response [51].

A study comparing anti-PD-1 monoclonal antibodies combined with lenvatinib or axitinib (a VEGFR-selective inhibitor) showed that both combination therapies exhibited higher anti-tumor activity and longer survival in mouse models compared to monotherapy. Furthermore, the anti-tumor activity of the anti-PD-1 monoclonal antibody combined with lenvatinib was enhanced compared to the combination with axitinib. Mechanistically, lenvatinib targeting FGFR was shown to reduce the number of tumor-associated macrophages and increase CD8 T cell counts, further highlighting its specificity and superior efficacy [48].

These preclinical data demonstrate the promising efficacy of lenvatinib combined with ICIs, which will facilitate the clinical translation of combination therapy. Further clinical trials are needed to validate the effectiveness and safety of this combination therapy in different liver cancer patient populations. With a deeper understanding of the mechanisms behind lenvatinib and ICI combination therapy, there is hope for optimizing treatment regimens to improve patient survival and quality of life. Ultimately, the successful implementation of this combination therapy could change the treatment standards for liver cancer and bring breakthroughs to clinical practice.

#### **Clinical efficacy of lenvatinib and ICIs in HCC**

##### **Comparative efficacy with sorafenib**

##### ***REFLECT trial: lenvatinib vs. sorafenib in advanced HCC***

Sorafenib is the standard first-line treatment for advanced liver cancer. Since its approval in 2007, multiple drugs have attempted to challenge Sorafenib's position but have failed. The REFLECT Trial [52] aimed to explore whether

Lenvatinib could be the first drug to match or even surpass Sorafenib. The study collected data from 954 treatment-naïve, unresectable advanced HCC patients across 154 centers in 20 countries from March 1, 2013, to July 30, 2015. Patients were randomly assigned in a 1:1 ratio to receive Lenvatinib ( $N=478$ ) or Sorafenib ( $N=476$ ). Participants took Lenvatinib (12 mg/d for those weighing  $\geq 60$  kg; 8 mg/d for those  $< 60$  kg) or Sorafenib (400 mg bid), with each cycle lasting 28 days. The median follow-up was 27.7 months for the Lenvatinib group and 27.2 months for the Sorafenib group. Results showed that for the primary endpoint, the median overall survival (OS) was 13.6 months for the Lenvatinib group and 12.3 months for the Sorafenib group, achieving non-inferiority (HR 0.92) but not superiority. However, all secondary endpoints (progression-free survival (PFS), time to progression (TTP), objective response rate (ORR), quality of life, and plasma pharmacokinetic exposure parameters) were significantly better in the Lenvatinib group. Additionally, Lenvatinib outperformed Sorafenib in all PFS subgroups. Both treatments had manageable safety profiles. This large, multicenter study demonstrated that Lenvatinib is not inferior to Sorafenib in overall survival for untreated advanced hepatocellular carcinoma, positioning Lenvatinib as a potential first-line standard treatment for advanced liver cancer.

#### **Other clinical trials: Lenvatinib vs. sorafenib in HCC**

Based on the REFLECT Trial, many subsequent studies have explored health-related quality of life (HRQOL) during treatment [53], subset analyses [54, 55], side effect studies [56, 57], and drug monitoring [58]. These clinical trials have laid a theoretical foundation for the broader application of Lenvatinib.

Results from two meta-analyses showed that the overall survival (OS), progression-free survival (PFS), objective response rate (ORR), and disease control rate (DCR) in the Lenvatinib monotherapy group were significantly higher than those in the Sorafenib group. However, the incidence of adverse events was higher in patients receiving Lenvatinib compared to those on Sorafenib [59], particularly regarding hypertension [60]. Additionally, studies indicated increased rates of hepatic encephalopathy [61] and renal impairment [62] in the Lenvatinib group. The incidence of hand-foot skin reactions was lower in the Lenvatinib group ( $P < 0.05$ ), while the rates of nausea, fatigue, urinary frequency, and dizziness showed no statistically significant difference between the two groups ( $P > 0.05$ ) [63]. For drug selection after HCC progression, one study evaluated the safety and efficacy of Lenvatinib versus Sorafenib in treating uHCC patients who progressed to Child–Pugh B (CP-B). The results indicated that Lenvatinib was more effective, with

manageable side effects, allowing patients with deteriorating liver function to continue Lenvatinib treatment [64]. Compared to Sorafenib, Lenvatinib may yield more favorable outcomes in advanced HCC with Vp3/4 [65]. Another study suggested that Lenvatinib could overcome Sorafenib resistance in HCC by intervening in the FGFR4-ERK signaling pathway, making it a suitable second-line treatment for unresectable HCC patients with Sorafenib resistance and FGFR4 expression [66].

Multiple studies have explored the benefits of Lenvatinib versus Sorafenib treatment. A cost-effectiveness analysis in China for patients with unresectable hepatocellular carcinoma showed that for patients weighing  $\geq 60$  kg, the incremental cost-effectiveness ratio (ICER) was close to an acceptable threshold, with Lenvatinib having a slightly higher probability of being cost-effective compared to Sorafenib. For patients weighing  $< 60$  kg, Lenvatinib demonstrated a comparative cost-effectiveness advantage [67]. The results of this study were more favorable than those from Japan [68]. In China, the direct medical cost and quality-adjusted life years (QALY) was 2.916, significantly higher than Japan's 1.88. A probabilistic sensitivity analysis from Canada indicated that in 64.87% of simulations, Lenvatinib remained a cost-saving option. Nevertheless, in the event of a 57% reduction in the cost of sorafenib, lenvatinib would cease to uphold its dominance as the primary strategy [69]. Similarly, a study from Australia also demonstrated that Lenvatinib is a cost-effective treatment choice compared to Sorafenib [70]. Through continuous assessments of treatment expenses and clinical results, Lenvatinib could emerge as the favored therapeutic choice for individuals afflicted with unresectable hepatocellular carcinoma. Future research should further investigate its long-term effects and adaptability in different patient populations to advance personalized treatment and provide better options for patients.

#### **Comparing the mechanism of action of lenvatinib with sorafenib**

Lenvatinib and Sorafenib are standard first-line treatments for HCC, both being TKIs, but they differ significantly in their mechanisms of action, primarily due to differences in targeted signaling pathways. Sorafenib targets the Raf/MEK/ERK pathway, as well as RTK, VEGFR1, VEGFR2, VEGFR3, and PDGFR- $\beta$ , while Lenvatinib targets VEGFR1–3, FGFR1–4, PDGFR- $\alpha$ , c-Kit, and RET, showing a notable difference in their kinase affinity profiles [71]. Both drugs share VEGFR as a common target; however, when VEGF and VEGFR are inhibited, FGF is activated to enhance FGFR signaling, promoting tumor angiogenesis. The activation of the FGF-FGFR signaling pathway serves as a compensatory

mechanism for tumor angiogenesis when VEGF-VEGFR pathways are suppressed. Compared to Sorafenib, Lenvatinib uniquely inhibits FGFR signaling and has a higher affinity for quickly binding to factors involved in tumor angiogenesis (such as VEGFR and FGFR), resulting in a strong inhibitory effect on tumor growth in vivo [30]. FGF21 may be a candidate biomarker for predicting Lenvatinib's potential to extend OS [72]. Additionally, Lenvatinib can reduce cancer stem-like cells (CSCs) in HCC by inhibiting FGFR1-3 signaling, a capability not possessed by Sorafenib [71].

Moreover, as a type V inhibitor, Lenvatinib binds to VEGFR2 more rapidly and with higher affinity, resulting in stronger kinase inhibitory activity compared to type II inhibitors like Sorafenib [72]. In cellular comparisons, while Lenvatinib is less effective than Sorafenib in inhibiting cell growth in vitro, it outperforms Sorafenib in inhibiting tumor growth in vivo [30]. Sorafenib is particularly active in highly differentiated cells with wild-type p53 and increased mitochondrial respiration. In contrast, Lenvatinib appears more effective in moderately to poorly differentiated cells with p53 mutations and lower mitochondrial respiration [73].

Understanding the different mechanisms and effects of these two drugs is crucial for developing personalized treatment plans. Continued research into the biological characteristics of HCC, combined with patient-specific factors, will aid in selecting the appropriate medication, thereby improving treatment outcomes and patient survival rates. Future clinical studies should focus on exploring the application potential of Lenvatinib in various types of liver cancer patients to support broader treatment strategies.

#### **Synergistic effects of lenvatinib and ICIs:**

##### **KEYNOTE-524: lenvatinib plus pembrolizumab**

The principle behind the combination of Lenvatinib and Pembrolizumab is that Lenvatinib can simultaneously inhibit tumor microenvironment-induced angiogenesis and immune suppression. This inhibition may enhance the clinical efficacy of PD-1 antibodies by boosting anti-tumor immune responses [74, 75]. KEYNOTE-524 is an open-label, single-arm Phase Ib clinical trial designed to explore the use of Lenvatinib and Pembrolizumab in unresectable HCC. The study included 104 patients with unresectable HCC, who received Lenvatinib (12 mg/day for patients weighing  $\geq 60$  kg; 8 mg/day for those  $< 60$  kg) in combination with Pembrolizumab (200 mg every 3 weeks). According to mRECIST and RECIST v1.1, the objective response rates (ORR) were 46.0% and 36.0%, respectively, with median durations of response of 8.6 and 12.6 months. The median progression-free survival (mPFS) was 9.3 and 8.6 months, while the median overall

survival (mOS) was 22 months. The results indicate that the combination therapy of Lenvatinib and Pembrolizumab yields a better response rate in unresectable HCC compared to Lenvatinib alone, with adverse events occurring within an acceptable range, demonstrating good anti-tumor activity and safety [76].

##### **LEAP-002: lenvatinib plus pembrolizumab vs. lenvatinib alone**

LEAP-002 [77] is a global, randomized, double-blind Phase III study aimed at assessing the efficacy and safety of Pembrolizumab in combination with Lenvatinib compared to Lenvatinib alone as a first-line treatment for advanced HCC. The results indicated that at the data cutoff, there were 252 deaths (64%) in the combination group and 282 deaths (71%) in the monotherapy group. The median overall survival (OS) was 21.2 months (95% CI: 19.0–23.6 months) for the combination group and 19.0 months (95% CI: 17.2–21.7 months) for the monotherapy group (HR: 0.84,  $p=0.023$ , not reaching the one-sided  $p=0.019$  superiority threshold), suggesting that the addition of Pembrolizumab did not significantly improve OS. In terms of progression-free survival (PFS), the median PFS was 8.2 months (95% CI: 6.4–8.4 months) for the combination group and 8.0 months (95% CI: 6.3–8.2 months) for the monotherapy group, indicating no significant difference in PFS between the two treatments. The combination of Pembrolizumab and Lenvatinib did not demonstrate the expected outcomes, with neither primary endpoint being achieved. Despite not reaching the primary endpoints, the combination therapy resulted in the longest OS observed to date for first-line treatment (21.2 months), while the OS for Lenvatinib monotherapy also exceeded expectations at 19 months. This clinical trial further supports the use of Lenvatinib monotherapy as a first-line treatment option for HCC patients.

##### **Other clinical trials combining lenvatinib and ICIs**

A study recruited adult patients with unresectable Barcelona Clinic Liver Cancer stage B or C HCC who were newly treated with systemic therapy. They received oral Lenvatinib in combination with intravenous anti-PD-1 agents (sintilimab/pembrolizumab/toripalimab/tislelizumab). The results showed a conversion success rate of 55.4% (31/56) for the primary endpoint. The objective response rate (ORR) according to mRECIST was 53.6% and 44.6% per RECIST 1.1. The median progression-free survival (PFS) was 8.9 months, and the median overall survival (OS) was 23.9 months. Among the 31 successfully converted patients, 21 underwent surgery, with an R0 resection rate of 85.7% and a pathological complete response rate of 38.1%. Grade  $\geq 3$  treatment-related adverse events were observed in 42.9%

of patients [78]. The results of the study on the efficacy and adverse reactions of Nivolumab combined with Lenvatinib for advanced HCC indicate that this combination can improve tumor control, reduce tumor burden, and enhance liver and immune function. Compared to Lenvatinib alone, the increase in adverse effects from the combination therapy was manageable [79, 80]. Additionally, clinical trials involving other ICIs [76, 81–84], including Tislelizumab combined with Lenvatinib, have shown good anti-tumor activity and tolerability, confirming the superiority of combination therapy.

In investigating treatment regimens, Wang et al. explored whether synchronous or asynchronous treatment with Lenvatinib and PD-1 inhibitors affected efficacy. The results indicated that the synchronous treatment group exhibited significantly improved OS and PFS compared to the asynchronous group, suggesting that administering Lenvatinib and PD-1 inhibitors together can significantly enhance survival in HCC patients [85]. Another study examined the impact of dosing and frequency on the efficacy of combination treatment, dividing patients into two groups: one receiving 6 mg/kg of Cadonilimab plus Lenvatinib every two weeks, and the other receiving 15 mg/kg every three weeks. Both groups achieved the expected treatment effects, but there were no significant differences in tumor control and safety between the two [86]. For patients excluded from the KEYNOTE-524 trial with tumor volume  $\geq 50\%$  of liver volume (TO  $\geq 50\%$ ) or Vp4 infiltration, Lenvatinib combined with Pembrolizumab may provide survival benefits for those with Vp4-infiltrating HCC, potentially expanding the indications for this combination therapy [87].

Compared to monotherapy, combination therapy has not shown significant statistical differences in safety across most clinical trials, but this does not mean that the side effects of combination therapy can be overlooked in clinical practice. Common adverse reactions include fatigue, loss of appetite, increased blood pressure, hand-foot skin reactions, diarrhea, rashes, acute asthma attacks, and muscle wasting, which should be managed during treatment [88, 89].

The broad application prospects of combination therapy have also led to the development of various predictive efficacy methods. Peripheral naive CD8<sup>+</sup> T cells, peripheral Th cells, and NK cells can predict responses to first-line combination therapy of Lenvatinib and anti-PD-1 antibodies in patients with advanced or unresectable HBV-related HCC [90, 91]. Using multivariate Firth logistic regression, Xu et al. confirmed that their proposed criteria are independent predictors for surgical conversion after Lenvatinib combined with anti-PD-1 antibody therapy [92]. Additionally, liver tumor stiffness measured by shear wave elastography [93], ALBI grading

[94], and MRI-based nomograms [95] have all been shown to predict the efficacy of Lenvatinib combined with PD-1 antibodies.

Lenvatinib in combination with immune checkpoint inhibitors (ICIs) urgently requires larger-scale clinical trials to validate its effectiveness and safety. These studies should encompass various types of hepatocellular carcinoma patients, considering factors such as age, sex, and disease duration to comprehensively assess treatment outcomes. Furthermore, exploring optimal dosing regimens and combination strategies will help improve patient survival and quality of life. Through systematic clinical research, we can better understand the mechanisms of this combination therapy, providing a solid theoretical foundation for clinical practice (Table 1).

#### **Multiple combination therapies based on ICI + lenvatinib**

In current research on hepatocellular carcinoma treatment, various combination therapies based on immune checkpoint inhibitors (ICIs) and Lenvatinib have garnered significant attention. This combinatorial approach aims to leverage the strengths of each agent, enhancing treatment efficacy and improving patient outcomes through synergistic effects. As exploration in this field deepens, an increasing number of studies are evaluating the potential applications of these combinations in diverse clinical contexts.

#### **ICI + lenvatinib + molecular targeted agent (MTA):**

In the treatment of hepatocellular carcinoma, exploring combination therapies for patients who experience progression after Lenvatinib monotherapy is particularly important. For these patients, the combination of immune checkpoint inhibitors (ICIs) with molecular targeted agents (MTAs) has demonstrated high anti-tumor activity and favorable safety profiles. Patients can receive ICIs and localized regional treatments while continuing Lenvatinib therapy, aiming for improved overall survival (OS) [96]. Research has also indicated that the failure to combine MTA adjunct therapy is considered an independent risk factor for tumor recurrence [97], further emphasizing the importance of comprehensive treatment strategies. Future studies should continue to focus on this area to optimize treatment regimens and enhance patient outcomes.

#### **ICI + lenvatinib + Hepatic artery infusion chemotherapy (HAIC):**

Compared to Lenvatinib alone, the combination of Lenvatinib, hepatic artery infusion chemotherapy (HAIC), and PD-1 inhibitors demonstrates a safe and promising anti-tumor activity in hepatocellular carcinoma (HCC) patients with high-risk features. The combination of

**Table 1** Clinical trials combining lenvatinib and ICIs

Clinical trial registration number	Nature of the test	Basis for clustering	Treatment target	ORR	PFS(months)	OS(months)	Safety and adverse reactions	Number of persons included	References
NCT03006926	Phase Ib multi-center, single-arm, open-label study	Lenvatinib + Pembrolizumab	Unresectable hepatocellular carcinoma	46%	9.3	22	TRAE Grade ≥ 3: 67%	104	[76]
ChiCTR1900023914	A single-arm, phase II trial	Lenvatinib + sintilimab/pembrolizumab/toripalimab/tislelizumab	Unresectable liver Cancer stage B or C HCC	53.60%	8.9	23.9	TRAE Grade ≥ 3: 42.9%	56	[78]
	Prospective clinical trial	Lenvatinib vs. lenvatinib + nivolumab	HCC	45.65% vs. 23.91%			NA	92	[79]
	A retrospective study	Lenvatinib + nivolumab vs. lenvatinib	Advanced hepatocellular carcinoma (aHCC)	45.0% vs. 23.4%	7.5 vs. 4.8	22.9 vs. 10.3	HR 0.2, 95% C.I. 0.1–0.7	87	[80]
NCT04401800	A multicenter, single-arm, phase 2 trial	Tislelizumab + lenvatinib	Unresectable hepatocellular carcinoma	38.70%	8.2	12: 88.6%	TRAE Grade ≥ 3: 28.1%	64	[81]
	A retrospective, unpaired, single-center study	Lenvatinib + anti-PD-1 antibodies	Advanced or unresectable hepatocellular carcinoma (HCC)	28.10%	8.4	17.2	AE 79.5%	210	[82]
NCT03892577	A retrospective study	Lenvatinib + anti-PD-1 antibodies	Unresectable hepatocellular carcinoma	19.60%	6.9	17.8	AE Grade ≥ 3: 57.9	378	[83]
	A retrospective study	Lenvatinib + anti-PD-1 antibodies vs. lenvatinib	Unresectable hepatocellular carcinoma	41.5% vs. 20.0%	8.0 vs. 3.0	not reached vs. 13.0	hypertension: 20% vs. 17.8%	127	[84]
	A prospective trial	Synchronous + asynchronous of lenvatinib + PD-1 inhibitor	HCC	35.9% vs. 19.8%	10.5 vs. 3.6	24.6 vs. 14.8	40.2% vs. 41.3%	213	[85]
NCT04444167	A phase Ib/II single-arm clinical trial	Time and dose of cardonizumab combined with lenvatinib	Advanced hepatocellular carcinoma (aHCC)	6 mg/kg/2 weeks: 35.5% 15 mg/kg/3 weeks: 35.7%	6 mg/kg/2 weeks: 8.6 months 15 mg/kg/3 weeks: 9.8 months	6 mg/kg/2 weeks: 27.1 months 15 mg/kg/3 weeks: NA	NA	59	[86]
	A retrospective study	Lenvatinib + pembrolizumab/toripalimab/sintilimab/nivolumab/camrelizumab	Tumor occupation ≥ 50% volume of liver (TO ≥ 50%) or invasion in Vp4	20.20%	6.6	11.4	NA	84	[87]

**Table 1** (continued)

Clinical trial registration number	Nature of the test	Basis for clustering	Treatment target	ORR	PFS(months)	OS(months)	Safety and adverse reactions	Number of persons included	References
NCT03892577	A multicenter observational retrospective real-world study	ICI+lenvatinib vs. ICI+others (apatinib/regorafenib/bevacizumab/sorafenib/donafenib)		13.8% vs. 3.7%	6.1 vs. 3.4	22.1 vs. 21.3	TRAE: 46.6% vs. 70.4%	85	[96]

hepatic artery infusion of FOLFOX with Lenvatinib and PD-1 inhibitors has proven effective as a first-line treatment, with manageable adverse effects [98–102]. It is noteworthy that HCC patients receiving HAIC combined with Lenvatinib and PD-1 inhibitors may experience HBV reactivation. Individuals experiencing HBV reactivation exhibit reduced survival durations in contrast to those without such reactivation. Consequently, it is advisable for patients with HBV-related hepatocellular carcinoma to undergo antiviral treatment and regular monitoring of HBV-DNA levels before and during combined therapeutic interventions [103]. Monitoring and managing HBV reactivation should be incorporated into clinical practice to ensure optimal treatment efficacy and patient outcomes. In the context of the Lenvatinib, HAIC, and PD-1 triplet therapy, levels of BTC (Betacellulin) and CCL28 (C–C motif chemokine ligand 28) may serve as predictive biomarkers for this combination treatment [104]. Additionally, AKR1C2+ and CFHR4+ hepatocyte subtypes may be predictive biomarkers for resistance to the combined therapy [105]. Future research should continue to focus on the exploration of biomarkers to optimize individualized treatment strategies.

#### **ICI + lenvatinib + transarterial chemoembolization (TACE):**

The combination of transarterial chemoembolization (TACE), Lenvatinib, and PD-1 inhibitors appears to significantly improve overall survival (OS), progression-free survival (PFS), objective response rate (ORR), and disease control rate (DCR) in patients with advanced hepatocellular carcinoma (HCC) without significantly increasing the risk of all-grade adverse events [106–119]. Nomograms can aid in stratifying treatment decisions and selecting appropriate populations for patients with unresectable HCC [120, 121]. Independent prognostic factors for survival include Child–Pugh classification, portal vein tumor thrombus (PVTT) classification, levels of IL-6, IL-17, IFN- $\alpha$ , and VEGF [122], as well as TAE scores [123]. The neutrophil-to-lymphocyte ratio and early tumor regression may serve as predictive biomarkers for patients with unresectable HCC [124]. These findings provide new insights for clinical treatment, suggesting that combinations of various therapeutic modalities may yield improved efficacy.

#### **Other combination therapies**

Furthermore, findings from clinical trials have indicated the tolerability and efficacy of Lenvatinib as a viable second-line systemic treatment after the administration of Atezolizumab and Bevacizumab in individuals diagnosed with advanced hepatocellular carcinoma (HCC) [125]. Furthermore, more complex combination therapies have shown promising clinical results, such as HAIC

combined with Lenvatinib and PD-1 antibodies. Concurrently, transarterial chemoembolization (TACE) and portal vein embolization (PVE) have been validated as safe and effective treatment options that can promote tumor necrosis in advanced HCC patients and enhance future liver volume [126]. The combination of TACE, Lenvatinib, PD-1 inhibitors, and I125 seed brachytherapy significantly improves overall survival (OS), progression-free survival (PFS), and disease control rate (DCR) in HCC patients with portal vein tumor thrombus (PVTT), demonstrating better prognosis [127].

In the treatment of liver cancer, the combined application of various therapeutic approaches is of great significance. This integrative treatment strategy maximizes the advantages of different modalities, addressing the complexity and heterogeneity of liver cancer to enhance treatment efficacy. By combining surgical interventions, pharmacotherapy (such as targeted drugs and immunotherapy), and localized treatments (like hepatic artery infusion chemotherapy and radiation therapy), more effective tumor control can be achieved while reducing the risk of recurrence and improving patient survival rates. Moreover, combination therapies can optimize individualized treatment plans, allowing for flexible adjustments based on disease progression, thus enhancing overall efficacy and quality of life. Therefore, exploring and implementing the combined use of multiple therapeutic strategies will be an important development direction in the field of liver cancer treatment (Table 2).

### **Future Perspectives of Lenvatinib in HCC Management**

#### **Biomarker-guided precision medicine:**

Biomarkers play a crucial role in the treatment of liver cancer with lenvatinib, as they can help predict patient responses and guide personalized treatment plans. Specific biomarkers are associated with treatment outcomes and can identify patients who are more likely to benefit from lenvatinib. Additionally, biomarkers can be used to monitor treatment responses and detect potential resistance mechanisms early, allowing for timely adjustments to the treatment regimen. Overall, integrating biomarkers into clinical practice enhances the precision of lenvatinib therapy, ultimately improving treatment outcomes and quality of life for liver cancer patients.

#### **Potential biomarkers for predicting response**

Specific gene expression in tumor tissues is significantly associated with the efficacy and survival benefits of lenvatinib. Research suggests that FGFR4 may predict the clinical benefits of lenvatinib for HCC patients [130]. KDM6A promotes HCC progression by activating FGFR4 expression, making it an important molecule



**Table 2** Multiple combination therapies based on ICI and Lenvatinib

Other treatments	Research type	Treatment type	ORR	PFS	OS	TRAE	Number of people	References
TACE + MWA + lenvatinib + anti-PD-1 antibodies vs. TACE + MWA	A retrospective study	HCC			21.257 months vs. 11.795 months	65.52%	67	[97]
HAIC-FOLFOX-lenvatinib-anti-PD-1 antibodies	A retrospective study	HCC with macrovascular invasion(MaVI)	31.20%	179 days	no reach	68.75%	32	[98]
HAIC-LEN-PD1 vs. LEN-PD1	A retrospective study	HCC patients exhibiting high-risk characteristics (Vp4, and/or bile duct invasion, and/or tumor occupancy ≥50%)	76.7% vs. 23.0%	9.6 months vs. 4.9 months	19.3 months vs. 9.8 months	higher	164	[99]
HAIC-FOLFOX-lenvatinib-anti-PD-1 antibodies	A retrospective study	uHCC(major portal vein tumor thrombosis (Vp3 and Vp4) or tumor occupancy exceeding 50% of the liver	52.7% (RECIST 1.1) 72.5% (mRECIST)	8.8 months	14.3 months	94.50%	91	[100]
HAIC-Len-ICI vs. Len-ICI	A retrospective study	Intermediate or advanced TACE-refractory HCC	48.30% vs. 23.80%	13 months vs. 7.2 months	24 months vs. 13 months	NA	121	[101]
HAIC(FOLFOX)-Len-PD1 vs. Len-PD1	A retrospective study	HCC patients with PVTT	61.8% vs. 20.8%	11.5 months vs. 5.5 months	26.3 months vs. 13.8 months	higher	142	[102]
HAIC-lenvatinib-anti-PD-1 antibodies	A retrospective study	HCC without HBV reactivation vs. CC with HBV reactivation	56.1% vs. 32.5%	6 months vs. 11.3 months	12.6 months vs. 25 months	lower	213	[103]
HAIC-lenvatinib-PD-1-inhibitors vs. HAIC-lenvatinib	a retrospective study	Advanced hepatocellular carcinoma refractory to hepatic arterial infusion chemotherapy			43.6 months vs. 18.9 months	NA	145	[117]
Lenvatinib, toripalimab plus FOLFOX-HAIC	phase II, single-centre, single-arm trial		63.90%	10.4 months	no reach		36	[104]
Lenvatinib, Toripalimab + FOLFOX		HCC with Extrahepatic Metastasis	43.30%	8.3 months	13.7 months		30	[105]
Pembrolizumab-lenvatinib-TACE vs. lenvatinib-TACE	A retrospective study	UHCC	47.1% vs. 27.8%	9.2 months vs. 5.5 months	18.1 months vs. 14.1 months	higher	142	[106]
PTL + TACE + Lenvatinib vs. TACE + Lenvatinib	A retrospective study	HCC patients with PVTT (portal vein tumor thrombus) types I-IV	NA	5.4 months vs. 2.7 months	NA	lower	41	[107]
TACE-lenvatinib-PD-1 inhibitor (TACE-LP) vs. TACE-TACE-L	A retrospective cohort study	AHCC		7.3 months vs. 4.0 months	16.9 months vs. 12.1 months		81	[108]

**Table 2** (continued)

Other treatments	Research type	Treatment type	ORR	PFS	OS	TRAE	Number of people	References
Lenvatinib-PD-1 inhibitor-TACE	A retrospective study	BCLC stage C HCC	mRECIST: 60% RECIST 1:1: 30%	8 months	18.4 months	93.30%	30	[112]
TACE-Lenvatinib and Camrelizumab vs. TACE+ Lenvatinib	A retrospective study	Unresectable Multiple Nodular and Large Hepatocellular Carcinoma (> 5 cm)	51.5% vs. 46.9%	9.4 months vs. 5.9 months	16.4 months vs. 11.0 months	33% vs 49%	82	[113]
TACE-lenvatinib -camrelizumab	A single-arm, open-label, multicenter, and prospective study	HCC with portal vein tumor thrombus	26.10%	9.3 months	18.2 months	TRAE ≥ 3: 34.8%	69	[114]
TACE-PD-1 inhibitors-lenvatinib vs. PD-1 inhibitors-lenvatinib	An observational, retrospective, cohort study	UHCC	54.3% vs. 25.4%	10.2 months vs. 7.4 months	20.5 months vs. 12.6 months	NA	105	[115]
TACE+ LEN+ PD-1 vs. LEN+ PD-1	A retrospective case-control study	UHCC	76.7 vs. 44.9%	16.2 months vs. 10.2 months	29.0 months vs. 17.8 months	NA	118	[116]
TACE- lenvatinib-PD-1 inhibitors vs. TACE+ len vs. TACE	A retrospective study	Unresectable recurrent HCC	70.4% vs. 48.9% vs. 42.5%	24.1 months vs. 17.3 months vs. 13.7 months	no reach vs. 25.6 months vs. 15.7 months	NA	204	[118]
PD-1 inhibitors, lenvatinib, TACE (PD-1-Lenv-T) vs. lenvatinib, TACE (Lenv-T)	A retrospective study	unresectable recurrent HCC	44.4% vs. 20%	11.7 months vs. 8.5 months	26.8 months vs. 14.0 months	NA	65	[128]
TACE-L-P vs. TACE-S (Sorafenib)-P	A retrospective study	HCC with Portal Vein Tumor Thrombus	41.25% vs. 30.59%	6.3 months vs. 3.2 months	21.7 months vs. 15.6 months	NA	165	[129]
TACE-lenvatinib (TACE-L)-PD-1 inhibitor vs (TACE-L)	A retrospective study	uHCC	54.0% vs. 32.8%	14.0 months vs. 9.0 months	24.0 months vs. 15.0 months	NA	241	[120]
TACE+ LEN+ PD-1 inhibitor vs. TACE+ LEN	A retrospective study	HCC with portal vein tumor thrombus	38.57% vs. 24.45%	7.5 months vs. 4.3 months	23.5 months vs. 18.3 months	NA	160	[122]
lenvatinib- atezolizumab -bevacizumab	A retrospective study	Advanced Hepatocellular Carcinoma	21.40%	4.2 months	8.3 months	100%	14	[125]
TACE-lenvatinib-PD-1 inhibitor, and iodine-125 seed brachytherapy vs. TACE+ len vs. TACE+ len+ PD-1	A retrospective study	HCC with portal vein tumor thrombosis		13 months vs. 5 months vs. 9 months	21 months vs. 10 months vs. 14 months	NA	150	[127]

influencing the efficacy of lenvatinib treatment in HCC; patients with high KDM6A levels are more likely to benefit from lenvatinib therapy [131]. Additionally, the expression of miR-3154 in tumor tissues may also predict the clinical benefits of lenvatinib for HCC patients [132].

At the cellular level, circulating tumor DNA (ctDNA) analysis stands out as a promising methodology offering significant clinical insights. Monitoring the dynamics of ctDNA within somatic cells of hepatocellular carcinoma (HCC) patients undergoing lenvatinib therapy emerges as a valuable biomarker for gauging disease progression. Frequently altered genes encompass TP53 (54%), CTNNB1 (42%), TERT (42%), ATM (25%), and ARID1A (13%). A reduction in the average variant allele frequency (VAF) following four weeks of lenvatinib treatment correlates with extended progression-free survival. While the mean mutation frequency during treatment bears prognostic relevance, it is important to note the absence of a discernible link between initial mutational status and the therapeutic efficacy of lenvatinib [133].

The physical function of patients is a crucial prognostic factor for the treatment of any disease, including hepatocellular carcinoma (HCC) treated with lenvatinib. In HCC patients receiving lenvatinib, skeletal muscle volume (SMV) [134], skeletal muscle index (SMI) [135, 136], geriatric nutritional risk index (GNRI) [137], fatigue status [138], and body composition assessments [139] are all significantly correlated with patient prognosis following treatment.

Imaging monitoring also plays a vital role in predicting the efficacy of lenvatinib treatment for HCC. The baseline volume index from 68 Ga-FAPI PET/CT is a potential independent prognostic factor for predicting durable clinical benefits, progression-free survival (PFS), and overall survival (OS) in uHCC patients receiving combined PD-1 and lenvatinib therapy [140]. Early modifications in the time-intensity curve (TIC) detected during the arterial phase of contrast-enhanced ultrasound (CEUS) represent promising early imaging markers for assessing the efficacy of lenvatinib [141]. Additionally, measuring changes in tumor blood flow via CEUS is one of the indicators for evaluating the effectiveness of lenvatinib in advanced HCC [142]. A reduction rate of CT attenuation values  $\geq 40\%$  is indicative of a favorable response to lenvatinib in highly malignant tumors [143]. Furthermore, CT values of necrosis post-lenvatinib treatment (N-CTav) and pre-treatment 18F-fluorodeoxyglucose PET scans can predict overall outcomes following lenvatinib therapy [144, 145]. Additionally, Bo et al. developed a valuable machine-learning radiomics model that demonstrates good performance in predicting the response of unresectable HCC to monotherapy with lenvatinib [146].

In addition to the biomarkers mentioned above, numerous studies have developed new grading systems or models to monitor the efficacy of lenvatinib. Examples include the PIMET score [147], albumin-bilirubin (ALBI) grade [148], modified ALBI (mALBI) grade [149], the corresponding albumin simplified (ALBS) grade [150], the glasgow prognostic score (GPS) is determined based on serum levels of C-reactive protein and albumin [151], the AFP-GGT-AGH scoring system [152], Child–Pugh classification, prognostic nutritional index (PNI) [153], lenvatinib prognostic index (LEP) [154], onodera's prognostic nutritional index (O-PNI) [155], CONUT score [156], and mG8 score [157]. Predictive models such as the "lenvatinib multivariable prognostic model" (MPML) [158] and the risk classifier of South China cohort (RCOSC) [159] demonstrate good discriminative ability, calibration, and applicability.

These biomarkers derived from tissue and cellular sources, along with imaging methods and the establishment of new evaluation grades or models, show strong performance in predicting lenvatinib efficacy. They can also identify patients who would benefit most from lenvatinib treatment, providing reliable tools for precision clinical decision-making in HCC patients. These biomarkers not only reflect the biological characteristics of tumors but also assist physicians in assessing patients' treatment tolerability and potential risks of adverse effects. Through personalized evaluations for different patients, medical teams can develop treatment plans that better align with individual circumstances, thereby enhancing treatment success rates and patients' quality of life. Moreover, regular monitoring of changes in these biomarkers allows for timely adjustments to treatment strategies, optimizing therapeutic outcomes and further advancing the development of personalized treatments for liver cancer.

#### **Liquid biopsy approaches**

Liquid biopsy is a technique that detects circulating tumor cells (CTCs), circulating tumor DNA (ctDNA), and circulating tumor RNA (ctRNA) through the collection of body fluid samples such as blood, urine, and saliva. Its non-invasive and highly repeatable nature provides comprehensive and personalized information in the diagnosis, treatment, and monitoring of tumors, thereby helping to improve patient survival rates and quality of life.

In the context of lenvatinib treatment for liver cancer, numerous studies have identified various circulating tumor biomarkers that can predict the efficacy of lenvatinib. Lenvatinib improves the immune microenvironment in HCC patients, preventing the exhaustion of effector cells and inhibiting the quantity and function of

immunosuppressive cells. Consequently, many immune markers can validate the efficacy of lenvatinib, including the CTL/Treg ratio [25], Th1 cells, and serum TNF- $\alpha$  levels [28]. The neutrophil-to-lymphocyte ratio (NLR) [160] also plays a significant role in efficacy prediction, while the proportion of eosinophils in the blood before and after treatment is not a viable marker [161]. Additionally, inflammatory indicators and their combination with other types of markers can effectively predict lenvatinib efficacy. For instance, C-reactive protein (CRP) levels [162], the C-reactive protein-to-albumin ratio (CAR) [163], and the platelet-to-lymphocyte ratio (PLR) [164] are all predictive of overall survival in unresectable HCC patients receiving lenvatinib.

Lenvatinib exerts its anti-cancer effects by targeting receptors, and high levels of VEGF, FGF21, and FGF19 in the blood often indicate poor patient outcomes. Other predictive indicators for HCC patients undergoing lenvatinib treatment include serum angiopoietin 2 (ANG2) [72, 165], M2BPGi [166], ST6GAL [167], changes in nitric oxide levels [168], alpha-fetoprotein, and the ratio of branched-chain amino acids to tyrosine (BTR) [169].

Genotypically, the GALNT14-rs9679162 “GG” genotype has been associated with longer overall survival in HBsAg-negative aHCC patients treated with lenvatinib [170]. The SNP combination patterns of NOS3 rs2070744 and FGFR4 rs351855 may serve as predictive factors for treatment response and prognosis in HCC patients receiving lenvatinib [171].

The importance of liquid biomarkers in lenvatinib treatment for liver cancer is increasingly recognized. These biomarkers can non-invasively reflect the dynamic changes in tumors, aiding early assessment of patient responses to lenvatinib and monitoring potential resistance mechanisms during treatment. Liquid biomarkers hold significant clinical application value in the treatment of liver cancer with lenvatinib, advancing the progress of personalized therapy for this disease (Table 3).

### Combination strategies:

#### *Lenvatinib plus other TKIs*

Combining lenvatinib with other tyrosine kinase inhibitors (TKIs) can integrate multiple target inhibitions, thereby addressing the biological characteristics of tumors more comprehensively, optimizing treatment strategies, and improving patient outcomes. Regorafenib, an oral broad-spectrum tyrosine kinase inhibitor, received FDA approval in April 2017 for use as a second-line therapy in patients displaying resistance to sorafenib. However, resistance to regorafenib also occurs. Research by Zhang et al. demonstrated that lenvatinib can reverse regorafenib resistance in hepatocellular carcinoma (HCC) by downregulating the IGF1R/

Mek/Erk signaling pathway, highlighting the potential of combining regorafenib with lenvatinib in HCC treatment [174]. Furthermore, prolonged exposure to sorafenib can lead to upregulation of FGFR1 in HCC cells, resulting in resistance to sorafenib, while lenvatinib may overcome this resistance by blocking FGFR1 [8]. Therefore, clinical studies on the combination of lenvatinib after sorafenib resistance are also anticipated.

#### *Lenvatinib plus anti-angiogenic agents*

One of the primary mechanisms by which lenvatinib exerts its anti-cancer effects is through the inhibition of tumor angiogenesis. When combined with other anti-angiogenic agents, it can enhance the overall anti-cancer efficacy. Concurrently, hepatocellular carcinoma (HCC) cells acquire resistance to lenvatinib through the activation of the epidermal growth factor receptor (EGFR) pathway, thereby inducing the EGFR-STAT3-ABCB1 axis. Targeting EGFR can help counteract the resistance induced by lenvatinib, as EGFR inhibition increases the sensitivity of liver cancer to lenvatinib. The combination of lenvatinib with epidermal growth factor receptor (EGFR) inhibitors represents a promising strategy that may improve clinical outcomes for certain patients [175]. Erlotinib, an inhibitor of the epidermal growth factor receptor (EGFR), has exhibited the capability to suppress ABCB1 activity, consequently diminishing the efflux of lenvatinib. Co-administration of lenvatinib with erlotinib has demonstrated substantial synergistic effects on hepatocellular carcinoma, as evidenced in both in vitro and in vivo studies [176]. Ramucirumab, a targeted agent that binds selectively to vascular endothelial growth factor receptor 2 (VEGFR-2) to hinder its activation, demonstrates promise in halting disease advancement among individuals with unresectable hepatocellular carcinoma (HCC) who have prior exposure to lenvatinib, all the while preserving hepatic function throughout treatment [177, 178]. Additionally, clinical trials have confirmed the efficacy and safety of the triple combination of lenvatinib with anti-angiogenic agents (such as bevacizumab) and PD-1 inhibitors (like atezolizumab) [179, 180].

The synergistic utilization of lenvatinib in conjunction with other tyrosine kinase inhibitors (TKIs) and anti-angiogenic agents displays auspicious clinical prospects. This strategy not only addresses numerous biological pathways implicated in tumor advancement but also proficiently retards the emergence of resistance mechanisms.

In addition, combinations of lenvatinib with proton beam therapy [181], hepatic arterial infusion chemotherapy [182–185], transarterial chemoembolization (TACE) [186], and lenvatinib-eluting microspheres [187, 188] have demonstrated promising clinical application potential. By integrating drugs with different mechanisms,

**Table 3** Biomarkers and other predictive tools

Predictor molecule	Location	Relationship with prognosis	References
FGFR4	Tissue	Positive correlation	[130]
miR-3154	Tissue	Negative correlation	[132]
Skeletal muscle volume (SMV)	Muscle	Positive correlation	[134–136]
KDM6A	Tissue	Positive correlation	[131]
Neutrophil-to-lymphocyte ratio	Blood	Negative correlation	[124]
ctDNA	Blood		[133]
CD8+T cell	Blood	Negative correlation	[90]
C-reactive protein to albumin ratio (CAR)	Blood	Negative correlation	[163]
Th cell, NK cell	Blood	Positive correlation	[91]
VEGF、FGF21 和 ANG2	Serum	Negative correlation	[72]
Mac-2-binding protein glycosylation isomer (M2BPGi)	Serum	Negative correlation	[166]
Platelet-lymphocyte ratio(PLR)	Blood	Negative correlation	[164]
CTL/Treg 比率	Serum	Positive correlation	[25]
GALNT14-rs9679162 genotype	Blood	Positive correlation	[170]
Eosinophil count	Blood	NA	[161]
ANG2 and VEGF	Serum	VEGF: Positive correlation ANG2: Negative correlation	[165]
TH1 cell, TNF- $\alpha$	Blood	Positive correlation	[28]
CRP	Serum	Negative correlation	[162]
ST6GAL	Serum	Positive correlation	[167]
NOS3 rs2070744 genotypes	Blood		[171]
2 M-DBR (DBR for the first 60 days), $\alpha$ -fetoprotein and branched-chain Amino acid to tyrosine ratio (BTR)	Blood	Positive correlation	[138, 169]
Reduction in tumor stain at 2 weeks	Blood, CT	Negative correlation	[172]
Predictive model/grading		Relationship with prognosis	References
A tumor-to-normal liver ratio (TLR)		Negative correlation	[145]
Cirrhotic PDX model			[173]
Modified albumin-bilirubin grade(mALBI)		Negative correlation	[149]
Onodera's prognostic nutritional index (O-PNI)		Negative correlation	[155]
The time-intensity curve (TIC) of arterial phase on contrast-enhanced ultrasound (CEUS)			[141]
The albumin-bilirubin (ALBI) grade		Negative correlation	[148]
Modified Geriatric 8 (mG8) score		Positive correlation	[157]
The Geriatric Nutritional Risk Index (GNRI)		Positive correlation	[137]
PIMET score		Positive correlation	[147]
Blood flow reduction		Positive correlation	[142]
MPML model		Negative correlation	[158]
CONUT score		Negative correlation	[156]
The albumin simplified (ALBS) grade		Negative correlation	[150]
Glasgow prognostic score (GPS)		Negative correlation	[151]
AFP-GGT-Hangzhou (AGH) scoring system		Negative correlation	[152]
Body composition parameters			[139]
Prognostic nutrition index (PNI)		Positive correlation	[153]
Machine learning (ML) radiomics models			[146]
Reductions in the rate of mean computed tomography (CT) attenuation values		Negative correlation	[143]
The lenvatinib prognostic index (LEP) index		Negative correlation	[154]
A Risk Classifier of South China Cohort (RCOSC)			[159]
The computed tomography (CT) attenuation value (CTav)		Positive correlation	[144]

**Positive correlation: the higher the content or value, the better the prognosis**

**Negative correlation: the lower the content or value, the better the prognosis**

these treatment regimens can more comprehensively address the complexities of liver cancer, thereby improving patient survival and quality of life. Future research will further validate the safety and efficacy of these combination therapies, providing more effective treatment options for liver cancer patients.

### Overcoming resistance to lenvatinib

#### *Mechanisms of resistance*

In the treatment of liver cancer, the emergence of resistance to lenvatinib limits its long-term efficacy. Data indicate that once patients develop resistance, the average duration of treatment is only 7.4 months [3]. Research indicates that resistance mechanisms are closely associated with factors such as genetic mutations in tumor cells, reprogramming of signaling pathways, and alterations in the tumor microenvironment. A deeper understanding of these resistance mechanisms is crucial for optimizing treatment strategies and improving patient outcomes.

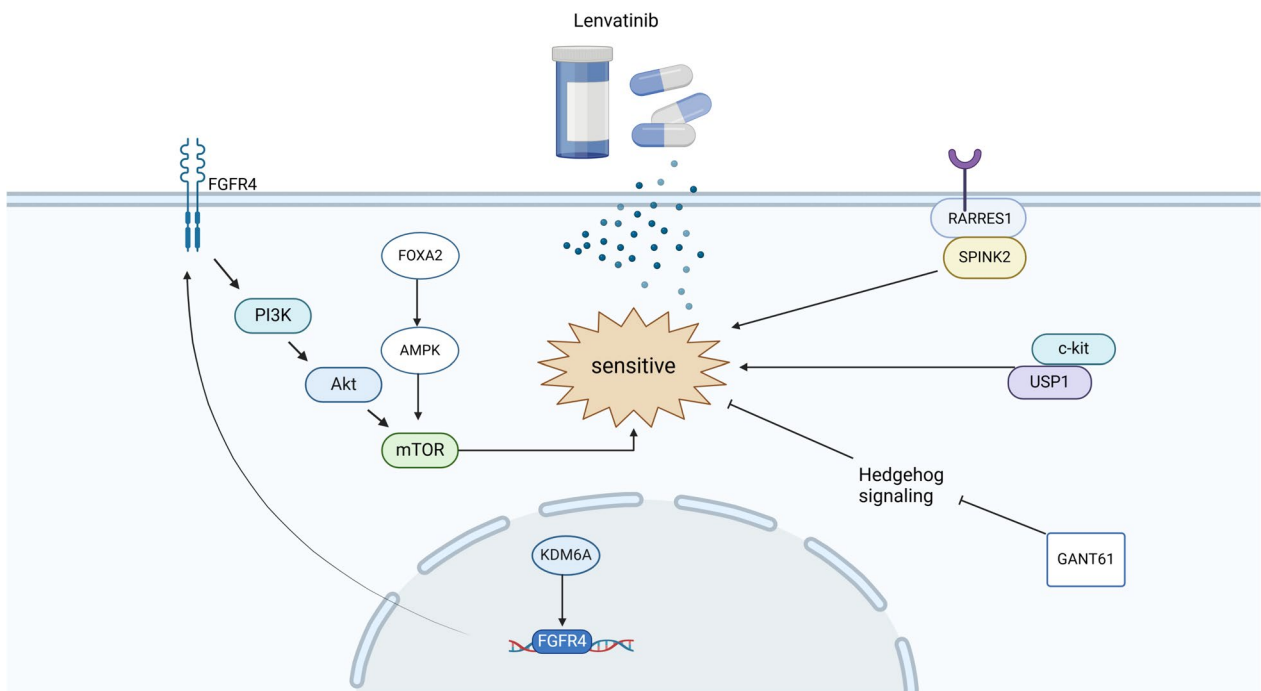
*Mechanisms for increasing sensitivity to lenvatinib* In the treatment of liver cancer, enhancing patient sensitivity to lenvatinib is crucial for improving efficacy. This can be explored from three aspects: coding genes, non-coding RNAs, and drug strategies. First, the regulation of certain coding genes can influence tumor cell responses to lenvatinib by optimizing target expression and signaling pathway activity. Second, non-coding RNAs, particularly specific miRNAs and long non-coding RNAs (lncRNAs) can promote drug sensitivity by modulating the expression of resistance-related genes. Finally, appropriate drug combination strategies and administration methods can significantly enhance the efficacy of lenvatinib through synergistic effects. By comprehensively studying these factors, new approaches for personalized treatment of liver cancer may be developed.

*Coding genes* in the context of coding genes, FOXA2 is a transcription factor essential for normal liver development and metabolism. Its overexpression in hepatocellular carcinoma (HCC) is often associated with better prognosis, as it enhances the inhibitory effect of lenvatinib on HCC cells by upregulating the AMPK-mTOR pathway [189]. Ubiquitin-specific protease 1 (USP1) is a key deubiquitinating enzyme that is overexpressed in HCC patients and plays a critical role in tumor progression. Mechanistic studies indicate that USP1 enhances the stability of the c-kit through interaction, making the c-kit an important target for lenvatinib treatment. Thus, the upregulation of USP1 sensitizes HCC cells to lenvatinib therapy [190]. GANT61, a selective inhibitor of GLI that suppresses the hedgehog signaling pathway, has demonstrated potential anti-cancer effects.

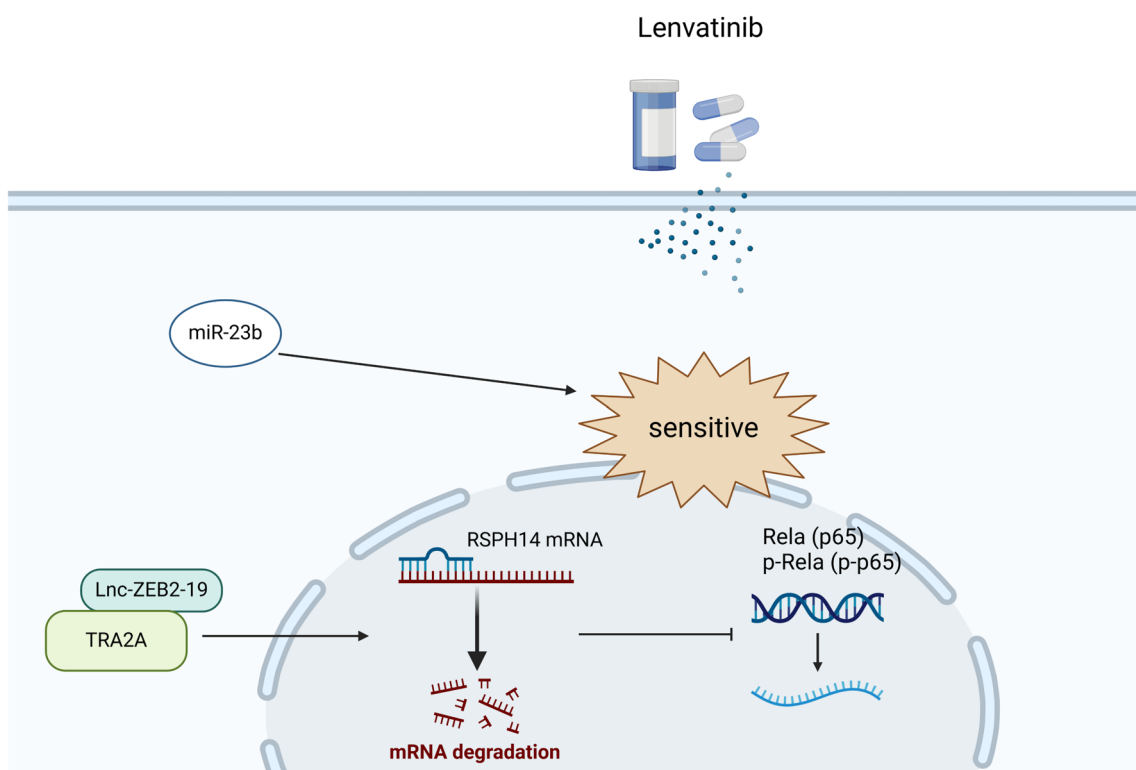
In HCC cells, particularly those with high CD133 expression, the combination of lenvatinib and GANT61 can reverse lenvatinib resistance and improve treatment efficacy [191]. RARRES1, a gene encoding a membrane protein, has been shown to interact with SPINK2 in HCC, promoting SPINK2 expression and inhibiting tumor cell proliferation and migration, thus increasing sensitivity to lenvatinib [192]. In HCC, KDM6A is significantly upregulated and associated with poor prognosis. KDM6A promotes FGFR4 expression, further activating the PI3K-AKT-mTOR signaling pathway, leading to metabolic reprogramming of glucose and lipids. Patients with high KDM6A expression in HCC exhibit greater sensitivity to lenvatinib-targeted therapy [131]. DUSP4 inhibits the lenvatinib resistance process by suppressing MAPK/ERK signaling. These mechanisms highlight the importance of specific coding genes in enhancing lenvatinib sensitivity in HCC treatment [193] (Fig. 2).

*Noncoding RNA* in the realm of non-coding RNA, miR-23b, as one of the most extensively studied miRNAs, plays a significant role in the development of many diseases, and hence is commonly targeted as a therapeutic intervention point. In the context of lenvatinib resistance in liver cancer research, it has been confirmed that miR-23b can inhibit the proliferation, migration, invasion, and angiogenesis of lenvatinib in human hepatocellular carcinoma cells, ultimately enhancing the sensitivity of lenvatinib [194]. In lenvatinib-resistant cell lines, elevated expression of Lnc-ZEB2-19, by specifically binding with TRA2A, leads to destabilization and subsequent degradation of RSPH14 mRNA, further resulting in the transcriptional downregulation of Rela (p65) and p-Rela (p-p65), inhibiting the stemness and migration of liver cancer, ultimately suppressing the progression and drug resistance of HCC [195] (Fig. 3).

*Drugs* in terms of pharmaceuticals, cisplatin, a common chemotherapy drug, induces phosphorylation of Chk1 and Chk2 by promoting ATM/ATR, further inhibiting cyclin B1 and Cdc2, preventing cell transition from G2 phase to M phase, thus inhibiting LR cell proliferation. Simultaneously, cisplatin is involved in the regulation of drug-resistant miRNAs (including 17 upregulated miRNAs and 6 downregulated miRNAs) [196]. Losartan, an angiotensin II receptor blocker, reduces lenvatinib-mediated VEGF-A and IL-8 production, directly inhibiting endothelial cell growth, thereby increasing sensitivity to lenvatinib-mediated vascular inhibition [197]. Metformin, besides its application in diabetes, also plays a role in anticancer activities. The combination of metformin and lenvatinib inhibits the AKT signaling pathway, thereby suppressing FOXO3 phosphorylation, leading to its nuclear accumulation, and synergistically inhibiting HCC growth and migration



**Fig. 2** Coding gene promotes lenvatinib sensitivity



**Fig. 3** Noncoding RNA promotes lenvatinib sensitivity

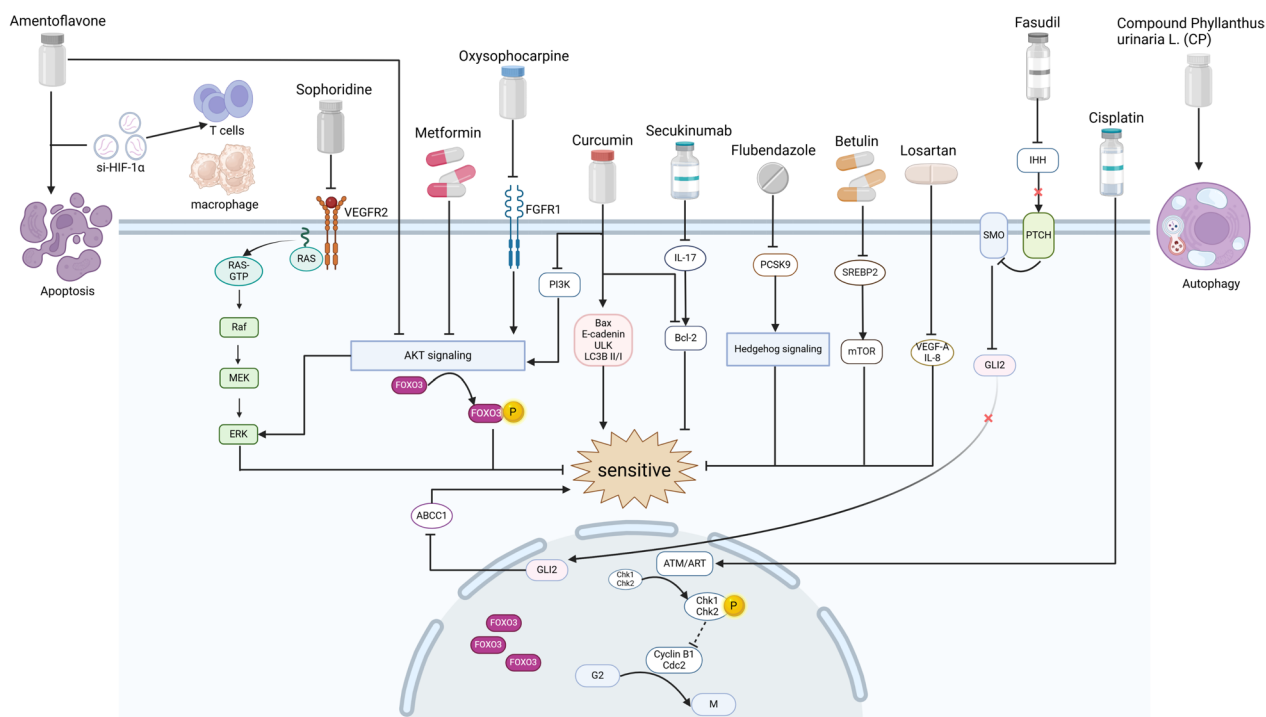
[198]. Secukinumab, the first approved IL-17A inhibitor globally, commonly used to treat psoriasis, when combined with lenvatinib, enhances tumor autophagy, better-suppressing tumor development compared to using lenvatinib alone [199]. Flubendazole, a common parasite treatment drug, has recently been reported for its role in cancer treatment, such as in breast cancer [200]. Its advantages in combination with lenvatinib have also been discovered in liver cancer [201]. Fasudil, a vasodilator, overcomes lenvatinib resistance in HCC by inhibiting the upstream switch IHH of the Hedgehog pathway, further inhibiting the GLI2-ABCC1 pathway activated by lenvatinib, ultimately overcoming lenvatinib resistance in HCC [202]. Traditional Chinese medicine also demonstrates great potential in enhancing sensitivity to lenvatinib. Betulinic acid, a natural pentacyclic triterpenoid compound widespread in birch trees and various plants, has shown promising prospects in anticancer applications and has recently emerged in enhancing the efficacy of lenvatinib. Betulinic acid inhibits SREBP2, further inhibiting the mTOR/IL-1 $\beta$  signaling pathway, demonstrating good sensitization effects both in vitro and in vivo. The combination therapy of lenvatinib and betulinic acid synergistically inhibits HCC cell proliferation [203]. Amentoflavone enhances sensitization effects by further enhancing lenvatinib's inhibition on the AKT/ERK signaling pathways and inducing apoptosis in liver cancer cells [204]. Sophoridine effectively inhibits tumor cell proliferation, particularly in LR cells, by reducing VEGFR2 expression and further suppressing the RAS/MEK/ERK axis to inhibit lenvatinib-resistant liver cancer cell growth [205]. Oxysophocarpine suppresses FGFR1 expression, subsequently inhibiting AKT and ERK signal transduction, making FGFR1-overexpressing HCC more sensitive to lenvatinib [206]. Compound *Phyllanthus urinaria* L. (CP), known for its potent anticancer effects, is extensively used in clinical treatment of liver cancer. The combination therapy of CP and lenvatinib effectively controls HCC progression by promoting exosome-mediated autophagy inhibition [207]. Curcumol effectively induces apoptosis in many cancer cells by targeting key signaling pathways such as MAPK/ERK, PI3K/Akt, and NF- $\kappa$ B. In LR cells, EGFR activation is significantly inhibited, and Curcumol enhances the efficacy of lenvatinib by inhibiting the EGFR-PI3K-AKT pathway [208]. Additionally, Curcumin has been reported to increase the expression of Bax, E-cadherin, ULK, and LC3B II/I while decreasing Bcl-2, N-cadherin, and JAK2/STAT3 expression, thereby enhancing lenvatinib's anti-tumor effects on liver cancer cells [209].

In conclusion, exploring strategies to enhance sensitivity to lenvatinib in liver cancer from the perspectives of coding genes, non-coding RNA, and pharmaceuticals

holds significant clinical importance. Regulating the expression of key genes, targeting specific miRNAs, and utilizing pharmaceutical agents in combination therapies are promising approaches to overcome lenvatinib resistance and improve treatment outcomes in hepatocellular carcinoma (Fig. 4).

*Mechanisms of increasing resistance to lenvatinib* Tyrosine kinase receptors lenvatinib exerts its anti-tumor effects by targeting tyrosine kinase receptors, including FGFR, EGFR, and MET receptors. Numerous studies have indicated that the activation or increased expression of these targeted receptors can promote the development of resistance to Lenvatinib. Lenvatinib inhibits FGFR4, further affecting GPX4 to induce ROS elevation, thereby mediating ferroptosis in tumor cells. Research has shown that Nrf2, a key regulator of antioxidant responses, can counteract Lenvatinib-induced ferroptosis by inhibiting ROS generation, and the inhibition of activated Nrf2 can resensitize HCC cells to Lenvatinib [14]. The upregulation of EGFR may lead to resistance development by promoting ROS accumulation [16, 210]. Additionally, studies have reported that the activation of the EGFR-STAT3-ABC1 axis significantly enhances Lenvatinib-mediated efflux-induced resistance, and the combination of Erlotinib, an EGFR inhibitor, with Lenvatinib has been shown to improve Lenvatinib resistance issues [176]. Moreover, it has been found that the two targets of Lenvatinib, EGFR, and FGFR, are not entirely independent, and their inhibition may lead to feedback activation of the EGFR-PAK2-ERK5 signaling axis, which is also a contributing factor to Lenvatinib resistance [16]. Activation of the hepatocyte growth factor (HGF)/c-MET signaling pathway is implicated in inducing resistance to Lenvatinib in hepatocellular carcinoma (HCC) cells exhibiting heightened c-MET expression levels. Combining Lenvatinib treatment with c-MET inhibitors can enhance systemic treatment efficacy in HCC patients [211]. Overexpression of ADAMTSL5 is associated with increased expression or phosphorylation of TKI receptors such as MET, EGFR, PDGFR $\beta$ , IGF1R $\beta$ , or FGFR4, thereby contributing to reduced sensitivity of HCC cells to TKIs including Lenvatinib [212]. METTL1 activates the EGFR pathway through m7G tRNA modification and participates in the development of Lenvatinib resistance in HCC patients [213]. Significantly increased METTL3 in Lenvatinib-resistant cells affects EGFR activation through m6A modification, further promoting Lenvatinib resistance [214]. The close relationship between the target receptors of Lenvatinib and resistance highlights the intricate nature of resistance mechanisms. In-depth exploration of these target receptors not only enhances our understanding of resistance mechanisms but also offers novel therapeutic





**Fig. 4** Drugs promotes lenvatinib sensitivity

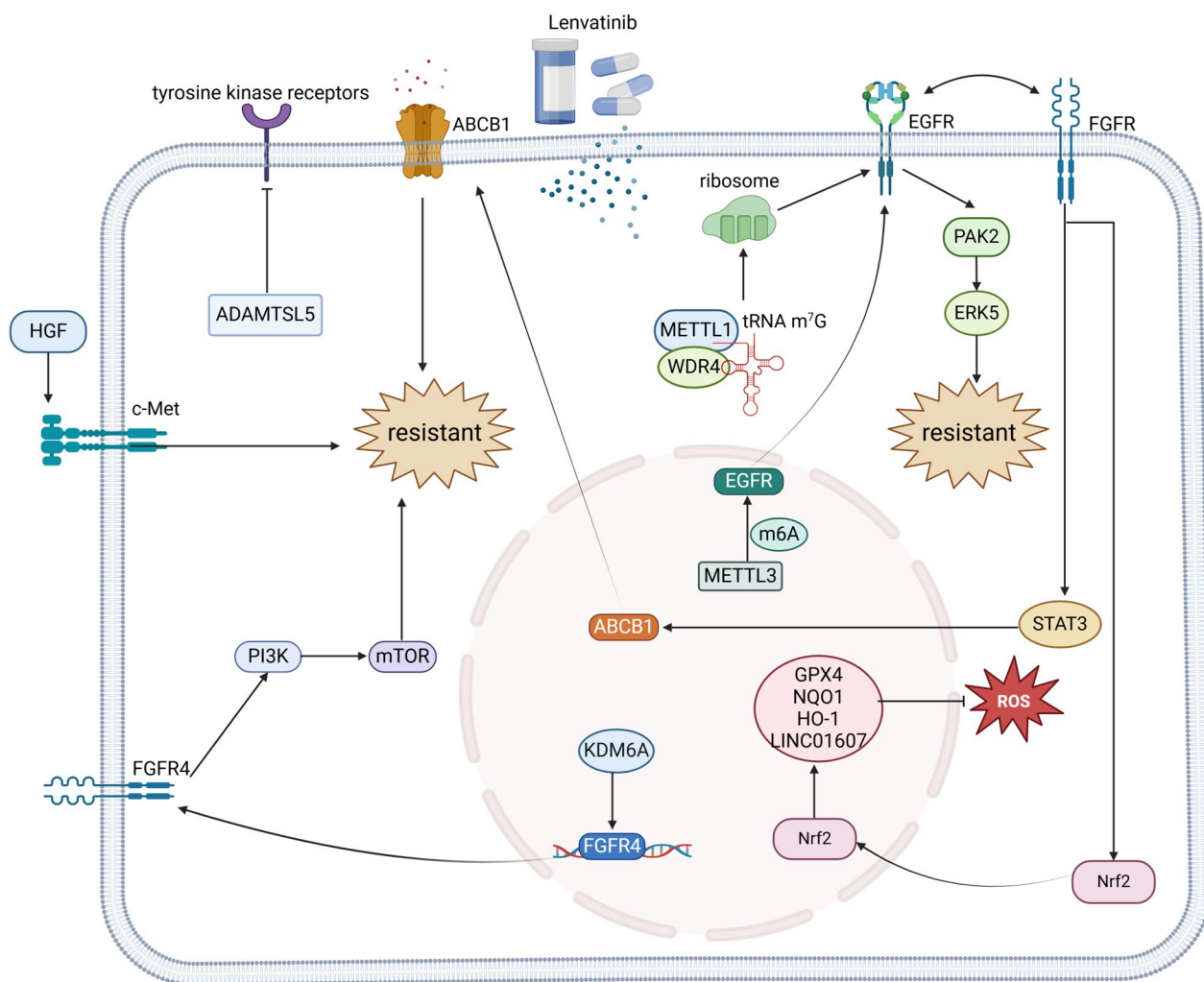
avenues to overcome resistance. By optimizing targeted treatment strategies and combining them with other therapies, there is potential to further enhance the efficacy of Lenvatinib, thereby improving treatment outcomes for liver cancer patients (Fig. 5).

*Signal pathways* in the treatment of liver cancer with lenvatinib, the role of signaling pathways is particularly crucial. These pathways not only regulate tumor cell growth and survival but also directly influence drug sensitivity and resistance. Investigating key signaling pathways can provide insights into how tumor cells develop resistance to lenvatinib. ITGB8, a member of the integrin beta chain family that encodes a type I membrane protein, promotes lenvatinib resistance by regulating AKT stability and enhancing AKT signaling through HSP90 [215]. The IGFBP-1-integrin  $\alpha 5\beta 1$  pathway facilitates tumor angiogenesis, contributing to HCC resistance [216]. In hypoxic environments, the PPARGC1A/BAMBI/ACSL5 axis is activated, regulating ROS production to inhibit ferroptosis in HCC cells [217]. Additionally, pathways such as TM4SF1-MYH9-NOTCH [218], FAK-WNK1 [219], USP22-JMJD8 [220], and METTL3-FZD10/ $\beta$ -catenin/c-Jun/MEK/ERK are all implicated in the development of lenvatinib resistance [221]. A deeper exploration of the significance of these signaling pathways in lenvatinib resistance will provide new insights and strategies for the personalized treatment of

liver cancer. The abnormal activation or reprogramming of these pathways is often a primary cause of resistance, highlighting their potential therapeutic value as targets for intervention (Fig. 6).

*Metabolic factor* in the treatment of liver cancer with lenvatinib, metabolic factors play a significant role and have a notable impact on the development of resistance. The metabolic reprogramming of tumor cells can alter their response to drugs, affecting drug uptake, metabolism, and excretion. Specific metabolic pathways, such as glycolysis, lipid metabolism, and amino acid metabolism, may play critical roles in resistance mechanisms. By studying these metabolic factors, we can gain a comprehensive understanding of how liver cancer cells adapt to lenvatinib treatment, providing new targets and strategies to overcome resistance.

During lenvatinib treatment, the drug can increase the recruitment of neutrophils by promoting the secretion of CXCL2 and CXCL5. Once in the tumor microenvironment (TME), neutrophils polarize to the N2 phenotype. Tumor-derived lactate can induce the expression of PD-L1 on neutrophils through the MCT1/NF- $\kappa$ B/COX-2 pathway. The increase in PD-L1-expressing neutrophils leads to the upregulation of Tregs and a decrease in T cell cytotoxicity. COX-2 inhibitors have been shown to reduce PD-L1-expressing neutrophils and restore T-cell cytotoxicity, which

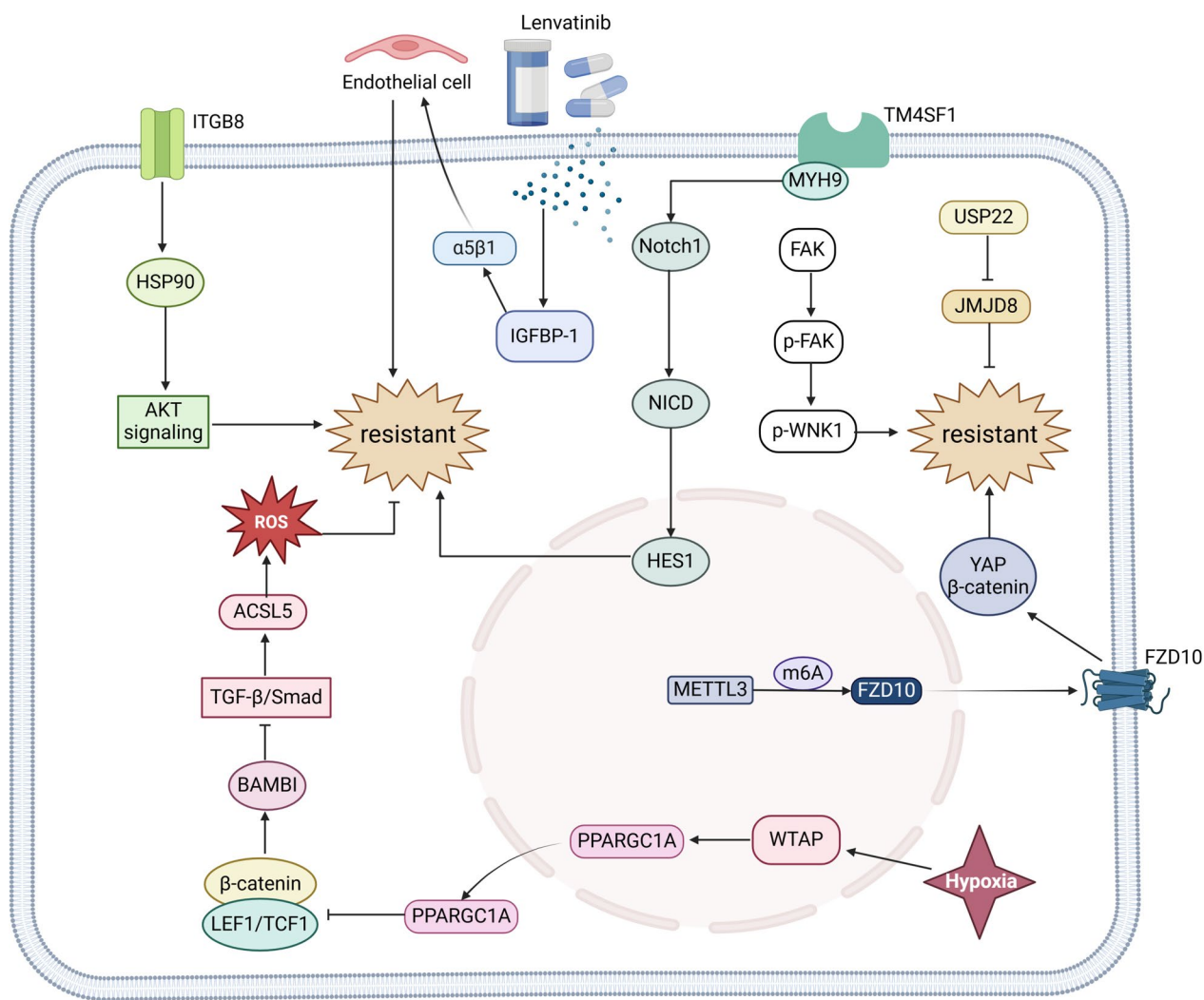


**Fig. 5** Targeting tyrosine kinase receptors promotes lenvatinib resistance

may provide a basis for improving lenvatinib resistance [222]. SLP2, a mitochondrial protein, can bind to JNK2, maintaining its stability. The increase in JNK2 enhances SREBP1 activity, promoting its translocation to the nucleus and facilitating de novo lipogenesis, thereby counteracting the effects of lenvatinib. This has also been validated in animal models [223]. ACYP1 can form a trimeric complex with HSP90 and MYC, promoting MYC nuclear entry and mediating the transcription of LDHA, which regulates glycolysis and promotes the progression of HCC. Targeting ACYP1 may synergistically enhance the effects of lenvatinib [224]. High expression of NQO1 reduces ROS generation induced by lenvatinib, thereby inhibiting apoptosis and promoting resistance to lenvatinib in liver cancer cells [225]. The significance of metabolic factors in resistance will provide important insights for the

personalization and optimization of liver cancer treatment (Fig. 7).

*Transcription factor* in the treatment of liver cancer with lenvatinib, transcription factors play a crucial role in regulating the expression of related genes, thereby influencing tumor cell responses and adaptations to the drug. Specific transcription factors may enhance the expression of resistance-related genes, leading to decreased sensitivity of tumor cells to lenvatinib. LEF1 has emerged as a new target to overcome lenvatinib resistance; studies have shown that in resistant cells, both LEF1 and its upstream regulator,  $\beta$ -catenin, are significantly elevated. This elevation affects the transcription of EMT-related genes (snail1, snail2, twist) in the nucleus, promoting resistance [226]. Under lenvatinib treatment, HIF-1 $\alpha$  levels increase in HCC cells, leading to enhanced transcription of STOML2. Increased STOML2 stabilizes PINK1, further enhancing mitochondrial autophagy, which

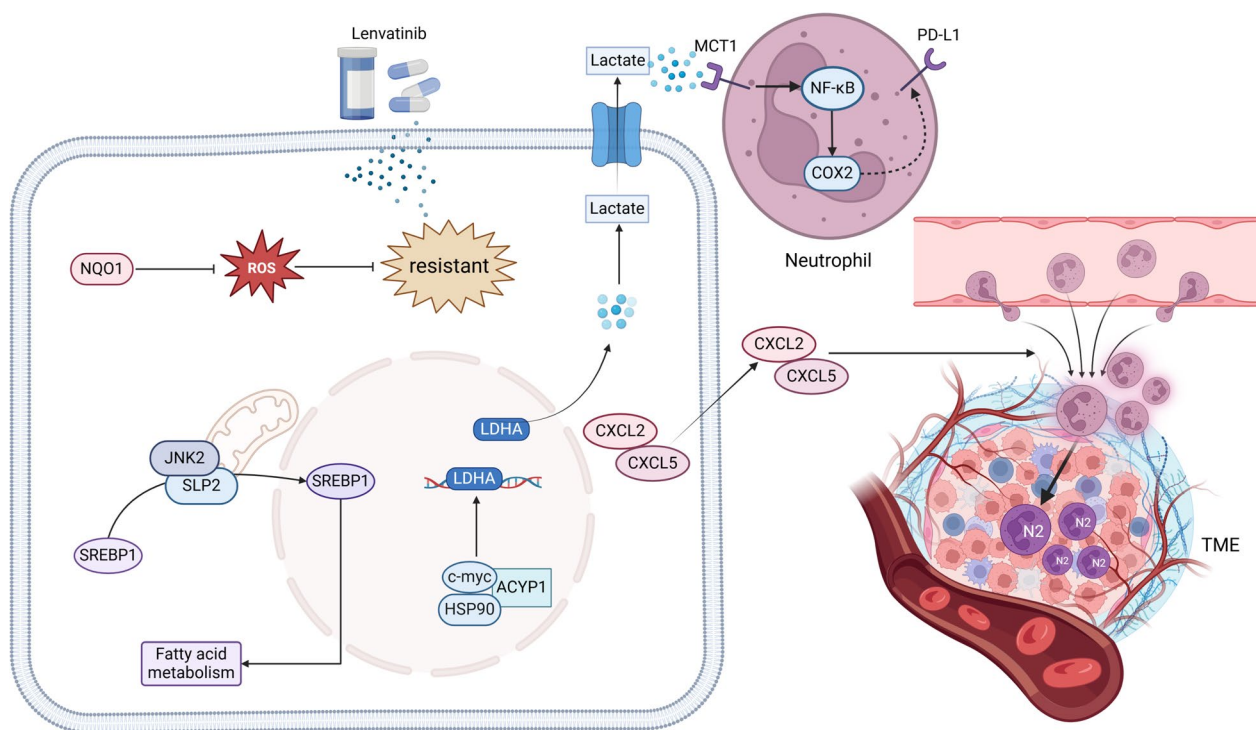


**Fig. 6** Promoting lenvatinib resistance through signaling pathways

supports normal physiological activities of tumor cells, promotes metastasis, and modulates HCC responses to lenvatinib [227]. HIF-1 $\alpha$  can also regulate autophagy levels by increasing the transcription of NRP1, contributing to the development of cellular resistance [228]. The DAGLA/2-AG axis promotes YAP nuclear translocation by inhibiting the phosphorylation of LATS1 and YAP, ultimately increasing the nuclear expression of PHLDA2, which can promote resistance [229]. WDR4 is highly expressed in liver cancer and contributes to lenvatinib resistance by promoting the expression of TRIM28, thus influencing downstream target gene expression [230]. The upregulation of CDK6 mediated by ERK/YAP signaling binds to and phosphorylates GSK3 $\beta$ , affecting LEF1 transcription and leading to the activation of Wnt/ $\beta$ -catenin, which in turn increases the expression of EMT-related genes, promoting lenvatinib resistance

[220]. YTHDF1 drives HCC resistance through the YTHDF1–m6A–NOTCH1 epitranscriptomic axis. Targeting YTHDF1 with lipid nanoparticles has been shown to significantly improve the efficacy of lenvatinib in HCC [231]. Investigating the mechanisms of these transcription factors not only helps unveil the fundamental causes of resistance but also provides potential targets for developing new therapeutic strategies (Fig. 8).

*Noncoding RNA* in the realm of non-coding RNAs, lnc MT1JP promotes resistance to lenvatinib treatment by targeting miR-24-3p, which elevates the levels of BCL2L2 and inhibits tumor cell apoptosis [232]. Studies have also shown that lncRNA MKLN1-AS plays a critical role in resisting lenvatinib-induced apoptosis [233]. Additionally, lncRNA AC026401.3 can bind to OCT1, facilitating its accumulation in the promoter region of E2F2, thereby activating the transcription of E2F2 and contributing to



**Fig. 7** Promoting lenvatinib resistance through metabolic factors

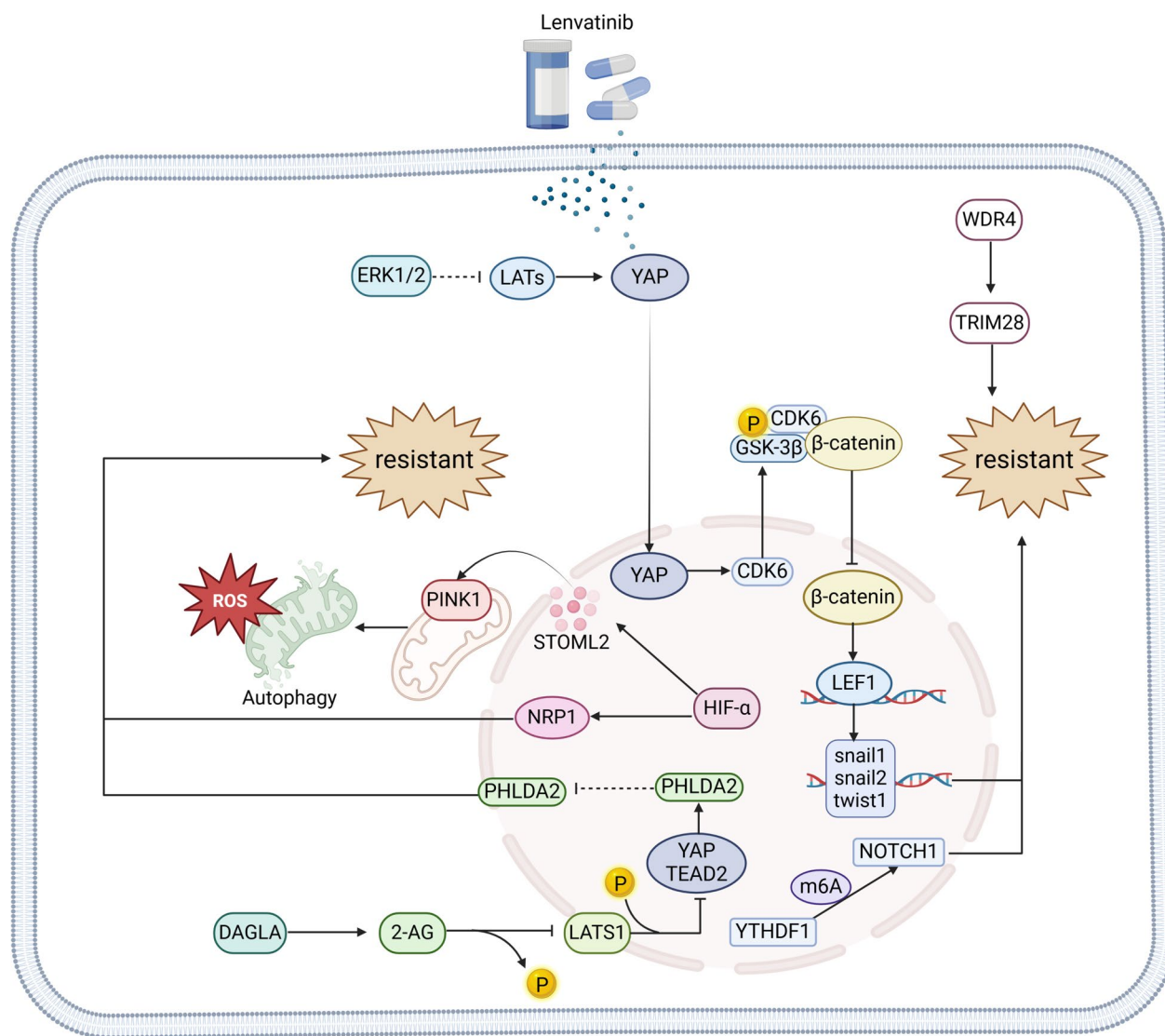
lenvatinib resistance in HCC [234]. LINC01607 competes with miRNA-892b, reducing its levels while upregulating P62. The overexpression of P62 can induce protective mitophagy and modulate the transcription of ferroptosis-related genes (such as GPX4 and HO-1) by promoting the nuclear entry of Nrf2, ultimately leading to decreased ROS production and enhanced lenvatinib resistance [235]. Lenvatinib-resistant cells can secrete circPAK1, which is transmitted to sensitive cells via exosomes, inducing resistance in those cells [236]. This underscores the critical role of ceRNAs in promoting lenvatinib resistance in HCC. Addressing the issue of lenvatinib resistance in liver cancer necessitates further exploration of ceRNA regulatory networks for targeted therapies (Fig. 9).

**Tumor immunity** in terms of immunity, SPP1 derived from cancer-associated fibroblasts (CAFs) influences the epithelial-mesenchymal transition (EMT) in liver cancer cells through the PKC $\alpha$ -RAF-MAPK-PI3K-AKT-MTOR pathway, leading to the development of resistance [237]. The combination of lenvatinib with immune checkpoint inhibitors has shown promising prospects compared to monotherapy; however, resistance issues still arise with combination treatments, and relevant studies have reported on this resistance. Zhou et al. found that lenvatinib combined with anti-PD1 antibody therapy leads to increased secretion of TNF by mucosal-associated invariant T (MAIT) cells in the immune

microenvironment. TNF acts on TNFRSF1B on Treg cells, which subsequently suppresses anti-tumor immunity [238] (Fig. 10).

**Drugs** in terms of pharmacology, targets that promote lenvatinib resistance have the potential as new therapeutic targets. Silencing or knocking out resistance-promoting genes has yielded promising results in cellular and preclinical studies, indicating potential for clinical translation. Notably, treatment strategies targeting HIF-1 $\alpha$  have attracted significant attention. In vitro, experiments have demonstrated that the use of attenuated *Salmonella* delivering siRNA-HIF-1 $\alpha$  in combination with lenvatinib effectively reduces the proliferation and angiogenesis of liver cancer cells while promoting apoptosis. Moreover, this combined therapeutic approach has the potential to augment the infiltration of T cells and M1 macrophages within the tumor microenvironment, amplifying the presence of immune cells in the spleen and thereby enhancing the host's anti-tumor immune response [239].

Predicting the occurrence of resistance in liver cancer patients during lenvatinib treatment is critically significant. Early identification of resistance risks allows for timely adjustments to treatment plans, improving therapeutic outcomes. Circulating tumor markers, due to their safety and sensitivity, are increasingly being utilized in cancer diagnosis and treatment. Analysis of circulating tumor DNA (ctDNA) in peripheral blood



**Fig. 8** Promoting lenvatinib resistance through transcription factor

provides a non-invasive method to identify cancer-specific mutations. Discovering heterogeneous genomic changes associated with lenvatinib resistance during treatment will greatly expand the application of ctDNA.

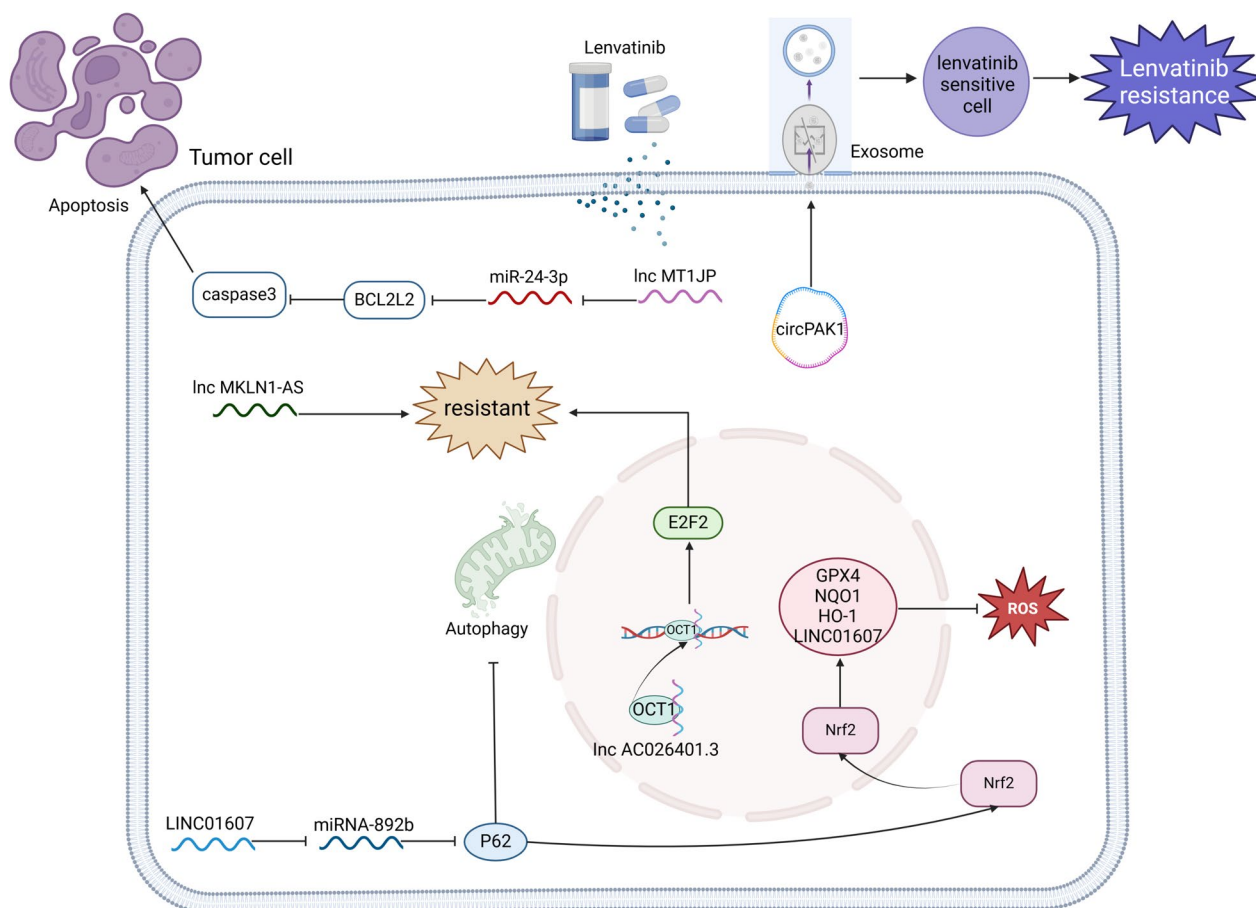
In summary, studying the mechanisms of lenvatinib resistance in liver cancer carries profound implications. This research not only helps elucidate the complex processes by which tumor cells adapt to drug pressure but also provides new insights and strategies for clinical treatment. By identifying and understanding these resistance mechanisms, a foundation can be established for developing new targeted therapies and combination treatment strategies, thereby improving patient outcomes and survival rates. Future research should

continue to focus on the diversity and complexity of resistance mechanisms to advance liver cancer treatment further.

**Emerging therapeutic modalities:**

**Linkage with nanotechnology**

The introduction of nanotechnology can enhance drug targeting and bioavailability, reduce side effects, and improve therapeutic efficacy. Lenvatinib, as a targeted therapy, can precisely target cancer cells and inhibit their growth. When combined with nanotechnology, the stability of the drug in the body is further improved, prolonging its circulation time while minimizing toxicity to normal cells, and offering new hope in cancer treatment.

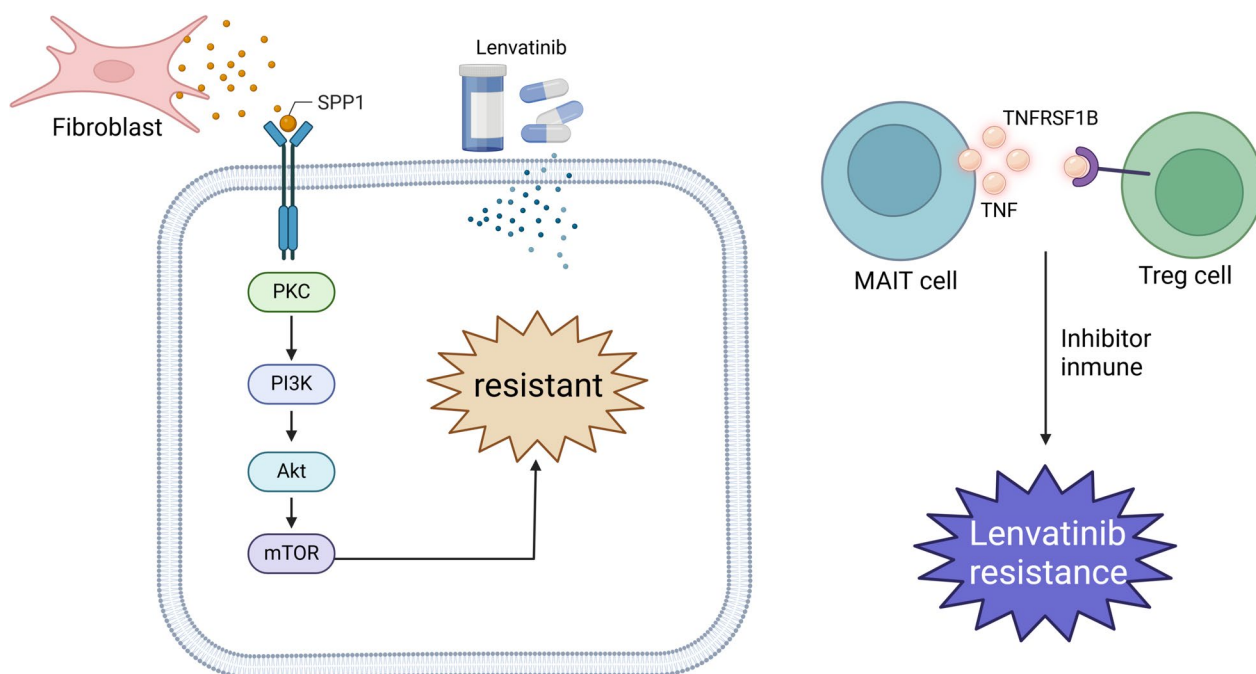


**Fig. 9** Promoting lenvatinib resistance through Noncoding RNA

Zhang et al. developed magnetic nanoliposomes (MNL) with dual-targeting capabilities for epithelial cell adhesion molecule (EpcAM) and wave protein, capable of encapsulating lenvatinib. In vitro and in vivo experiments demonstrated low cytotoxicity, effective inhibition of HCC cell proliferation, promotion of HCC cell apoptosis, and specific targeting capabilities with MRI tracing ability [240]. Wu et al. successfully constructed a biomimetic nanodrug delivery platform with lenvatinib at its core, wrapped in a pH-sensitive polymer, poly( $\beta$ -amino ester)-polyethylene glycol amine (PAE-PEG-NH<sub>2</sub>), and coated with cancer cell membrane (CCM). In vivo experiments indicated that this nanodrug exhibited superior therapeutic effects in subcutaneous tumor mouse models, effectively eliminating tumors [241]. Liu et al. developed Bi/Se nanoparticles (NPs) loaded with lenvatinib (Bi/Se-Len NPs), which demonstrated excellent performance in reversing hypoxic and immunosuppressive states in HCC. Additionally, these nanoparticles can be used for CT

image-guided stereotactic body radiotherapy (SBRT) in mice, providing accurate imaging guidance [242].

Metal-based nano drugs show promising clinical translational potential when combined with lenvatinib. Zeng et al. developed a targeted therapeutic strategy that co-assembled lenvatinib, adriamycin, and Fe<sup>3+</sup>, along with metal nanodrugs to enhance tumor vascular normalization and increase the infiltration of T lymphocytes within tumors. This approach improves anti-tumor immunity mediated by calreticulin by alleviating hypoxia, reducing regulatory T cells, and downregulating tumor PD-L1 expression, effectively inhibiting the progression of HCC in both in situ models and patient-derived organoids [243]. Another metal nanodrug, CAL@PG, was successfully constructed by Xu et al. In this technique, ultrasmall gold nanoparticles (AuNPs) and ultrafine copper sulfide nanocrystals (Cu<sub>2</sub>-xS NCs) were uniformly encapsulated within galactosamine-conjugated poly(lactide-co-glycolide) (PLGA) to form drug delivery nanoparticles. CAL@PG



**Fig. 10** Promoting lenvatinib resistance through tumor immunity

nanoparticles exhibited excellent stability under physiological conditions, while rapidly releasing lenvatinib in the unique tumor microenvironment (TME) and at elevated temperatures. This combination therapy significantly enhanced the efficacy of lenvatinib through near-infrared II (NIR-II) photothermal effects [244, 245]. Luo et al. loaded lenvatinib onto mesoporous Fe<sub>3</sub>O<sub>4</sub> (mFe) nanoparticles, with bovine serum albumin (BSA) attached to the NP surface to create a metal drug (BSA-mFe@Len NPs). This metal drug induces immune modulation in the TME against HCC through multiple functions of mFe NPs and lenvatinib, including triggering tumor apoptosis, vascular normalization, recruitment of cytotoxic T lymphocytes, and elimination of regulatory T cells. In vivo experiments demonstrated that this metal drug significantly inhibited hepatocellular carcinoma growth and evoked long-term anti-tumor immune memory, paving a new pathway for treating advanced HCC patients [246].

The combination with nanotechnology provides a multifunctional delivery system for lenvatinib, enriching the multimodal synergistic treatment approach for HCC. This method effectively enhances survival rates and quality of life for cancer patients while introducing new ideas and methods in cancer treatment. However, despite the potential shown in laboratory and clinical trials, further research and validation are needed to ensure safety and efficacy. As the fields of nanotechnology and biomedicine continue to advance, the

application prospects of combining nanotechnology with targeted drugs like lenvatinib will become even broader.

**Autologous platelets encapsulated the drugs**

Hepatocellular carcinoma activates platelets, which then bind to tumor-associated endothelial cells and release growth factors, promoting tumor progression. Hiroki et al. utilized platelets as drug carriers for tumor therapy. In both in vitro and in vivo experiments, autologous platelets encapsulating various tyrosine kinase inhibitors (lenvatinib or sorafenib) were shown to effectively deliver anti-cancer drugs to tumor tissues. The use of autologous platelets for drug encapsulation may represent a novel therapeutic strategy for HCC [247].

**Combination with Histone Deacetylase Inhibitors**

Histone deacetylase inhibitors (HDAC inhibitors) alter chromatin structure and gene expression by inhibiting the activity of histone deacetylases, leading to increased histone acetylation and enhanced gene transcription. This process affects cell proliferation, differentiation, and apoptosis. HDAC inhibitors show significant potential in cancer therapy and can enhance the effects of other treatments. In hepatocellular carcinoma, the combination of HDAC inhibitors with lenvatinib has demonstrated synergistic anti-tumor effects. Research by Ryo et al. indicated that class IIa inhibitors combined with lenvatinib induce apoptosis by downregulating FGFR4

and blocking FGFR signaling in FGFR4-positive HCC cell lines [248]. Moreover, Jane et al. discovered that the synergistic action of entinostat, an HDAC inhibitor, in conjunction with lenvatinib induces cell death by triggering ROS-dependent ATM activation and eIF2 $\alpha$  inactivation. Consequently, this series of events results in heightened cytotoxic autophagosome generation and reduced levels of protective mitochondrial protein expression [249].

## Discussion and conclusion

Lenvatinib is a multi-targeted tyrosine kinase inhibitor that exerts its anti-tumor effects primarily by inhibiting vascular endothelial growth factor receptors (VEGFR), fibroblast growth factor receptors (FGFR), and other related signaling pathways. By blocking these pathways, lenvatinib effectively suppresses tumor cell proliferation and survival, inhibits tumor angiogenesis, and reduces tumor blood supply, thereby achieving therapeutic efficacy. Additionally, lenvatinib has shown improvements in the liver cancer microenvironment, capable of altering the tumor-associated immune status and promoting anti-tumor immune responses.

In multiple clinical trials, lenvatinib has demonstrated good anti-tumor activity and a high objective response rate, particularly among patients with advanced hepatocellular carcinoma (HCC), where its efficacy has been widely recognized. Compared to traditional treatment methods, lenvatinib not only prolongs patient survival but also enhances quality of life. However, its clinical application is not without side effects, such as hypertension, fatigue, and liver function impairment. These adverse reactions may affect patient compliance and overall survival rates. Therefore, in clinical practice, it is essential to strengthen the monitoring and management of these side effects to improve the patient treatment experience.

Although Lenvatinib demonstrates promising efficacy in the treatment of liver cancer, the issue of drug resistance is becoming increasingly severe. Studies indicate that HCC cells may develop resistance through various mechanisms, including genetic mutations, signaling pathway alterations, and changes in the tumor microenvironment. The emergence of these resistance mechanisms not only impacts the effectiveness of individual drugs but may also lead to disease progression. Understanding the complexity of drug resistance mechanisms is crucial for the development of new therapeutic strategies. Research into resistance mechanisms can provide novel biomarkers for clinical use. Biomarkers play a significant role in early diagnosis, prognosis assessment, and treatment monitoring of liver cancer. By identifying resistance-related biomarkers, it becomes possible to better predict patient responses and tailor treatment plans.

Additionally, biomarkers can aid in identifying patient populations suitable for combination therapies, thereby enhancing overall treatment outcomes.

Combination therapies not only enhance anti-tumor immune responses but may also overcome resistance issues associated with single treatments. However, combining immune checkpoint inhibitors with Lenvatinib presents challenges in terms of safety and tolerability. Studies suggest that immune checkpoint inhibitors may trigger immune-related adverse events such as rash, hepatitis, and endocrine disorders, while Lenvatinib itself can lead to issues like hypertension and abnormal liver function. When these two drugs are used together, the safety risks for patients may significantly increase. Therefore, close monitoring of patient health status is essential during combination therapy to promptly identify and manage potential side effects, ensuring treatment safety and efficacy. Furthermore, further research should focus on the long-term safety of such combination therapies to provide more reliable treatment options for patients.

In conclusion, Lenvatinib holds significant clinical value in the treatment of liver cancer. Given the advantages of combination therapy, future research is anticipated to explore emerging treatment modalities involving Lenvatinib, such as combinations with novel ICIs (anti-LAG-3), CAR-T cell therapy, oncolytic viruses, ablation techniques, and other drugs or technologies. Understanding Lenvatinib resistance mechanisms, identifying biomarkers, and optimizing combination treatment strategies with immune checkpoint inhibitors offer new possibilities for improving the survival rates and quality of life for liver cancer patients. Future research should continue to focus on these areas to drive further advancements in liver cancer treatment.

## Abbreviations

ADCC	Antibody-dependent cell-mediated cytotoxicity
ALBI	Albumin-bilirubin
ALBS	Albumin simplified
ANG2	Angiopoietin 2
APCs	Antigen-presenting cells
AuNPs	Ultrasmall gold nanoparticles
Bi/Se-Len NPs	Bi/Se nanoparticles loaded with lenvatinib
BSA	Bovine serum albumin
BTR	Branched-chain amino acids to tyrosine
CAFs	Cancer-associated fibroblasts
CAR	C-reactive protein to albumin ratio
CAR-T	Chimeric antigen receptor T-cell
CCM	Cancer cell membrane
CCL28	C-C motif chemokine ligand 28
CEUS	Contrast-enhanced ultrasound
CP	Compound <i>Phyllanthus urinaria</i> L.
CP-B	Child-Pugh B
CRP	C-reactive protein
CSCs	Stem-like cells
CTav	CT values of necrosis post-lenvatinib treatment
CTLA-4	Cytotoxic T lymphocyte-associated protein 4
CTLs	Cytotoxic T lymphocytes
CTCs	Circulating tumor cells



CTD	Circulating tumor DNA
CTIC	Time-intensity curve
CTIM-3	T cell immunoglobulin and mucin domain-containing protein 3
CTIGIT	T cell immunoglobulin and ITIM domain protein
Cu <sub>2</sub> -xS NCS	Ultrafine copper sulfide nanocrystals
DCR	Disease control rate
DOR	Duration of response
ECM	Extracellular matrix
EMT	Epithelial-mesenchymal transition
EpCAM	Epithelial cell adhesion molecule
ERK	Extracellular regulated protein kinase
FGL1	Fibrinogen-like protein 1
FGF	Fibroblast growth factor
FGFR	Fibroblast growth factor receptors
FOLFOX	Oxaliplatin, 5-fluorouracil, and leucovorin
Gal-9	Galectin-9
GNRI	Geriatric nutritional risk index
GPS	Glasgow Prognostic Score
HAIC	Hepatic artery infusion chemotherapy
HCC	Hepatocellular carcinoma
HDAC inhibitors	Histone deacetylase inhibitors
HRQOL	Health-related quality of life
ICB	Immune checkpoint blockade
ICER	Incremental cost-effectiveness ratio
ICIs	Immune checkpoint inhibitors
KIT	KIT proto-oncogene receptor tyrosine kinase
LAG-3	Lymphocyte activation gene 3
LEP	Lenvatinib Prognostic Index
lncRNAs	Long non-coding RNAs
MAIT	Mucosal-associated invariant T
MDSCs	Myeloid-derived suppressor cells
mALBI	Modified ALBI
MHC	Major histocompatibility complex
MNL	Magnetic nanoliposomes
mOS	Median overall survival
MPML	Lenvatinib multivariable prognostic model
MSI-H	Microsatellite instability
mFe	Fe3O4
mPFS	Median progression-free survival
MTA	Molecular targeted agent
NCTAv	CT values of necrosis post-lenvatinib treatment
NIR-II	Near-infrared II
NAD <sup>+</sup>	Nicotinamide adenine dinucleotide
NLR	Neutrophil-to-lymphocyte ratio
OS	Overall survival
O-PNI	Onodera's Prognostic Nutritional Index
ORR	Overall response rate
PAE-PEG-NH <sub>2</sub>	Poly( $\beta$ -amino ester)-polyethylene glycol amine
PD-1	Programmed cell death protein 1
PD-L1	Programmed cell death ligand 1
PDX	Patient-derived xenograft
PDGF	Platelet-derived growth factor
PDGFR $\alpha$	Platelet-derived growth factor receptor alpha
PFS	Progression-free survival
PLGA	Galactosamine-conjugated poly(lactide-co-glycolide)
PLR	Platelet-to-lymphocyte ratio
PKB/AKT	Protein kinase B
PNI	Prognostic nutritional index
PSGL-1	P-selectin glycoprotein ligand 1
PVE	Portal vein embolization
PVTT	Portal vein tumor thrombus
QALY	Quality-adjusted life years
RCOSC	Risk classifier of south China cohort
RET	Rearranged during transfection
SBRT	Stereotactic body radiotherapy
SHP-2	Src homology region 2-containing protein tyrosine phosphatase 2
SIRP $\alpha$	Signal regulatory protein $\alpha$
SMI	Skeletal muscle index
SMV	Skeletal muscle volume

Syk	Spleen tyrosine kinase
STING	Stimulator of interferon genes
TACE	Transarterial chemoembolization
TCR	T cell receptor
TBV	Tumor blood volume
TCTDNA	Circulating tumor DNA
TCTRNA	Circulating tumor RNA
Th	T helper
TIGIT	T cell immunoglobulin and ITIM domain protein
TIME	Tumor immune microenvironment
TIM-3	T cell immunoglobulin and mucin domain-containing protein 3
TILs	Tumor-infiltrating lymphocytes
TME	Tumor microenvironment
TNPs	Bi/Se nanoparticles loaded with lenvatinib
Tregs	Regulatory T cells
TTP	Time to progression
TRAE	Treatment-related adverse events
USP1	Ubiquitin-specific protease 1
VAF	Variant allele frequency
VEGF	Vascular endothelial growth factor
VEGFR	Vascular endothelial growth factor receptors
VISTA	V-domain Ig suppressor of T-cell activation
VSIG3	V-set and immunoglobulin domain containing 3

#### Author contributions

YC and SD retrieved the data and drafted the manuscript. CC retrieved the data and revised the manuscript. LC initiated the idea and drafted the manuscript.

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#### Data availability

No datasets were generated or analysed during the current study.

#### Declarations

#### Competing interests

The authors declare no competing interests.

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