

The progress and challenge of anti-PD-1/PD-L1 immunotherapy in treating non-small cell lung cancer

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Abstract: Abstract

The use of programmed cell-death protein 1 (PD-1)/programmed cell-death ligand 1 (PD-L1) inhibitors is the standard therapy for the first-line or second-line treatment of patients with non-small-cell lung cancer (NSCLC). In contrast to current traditional treatments such as chemotherapy or radiotherapy, anti-PD-1 and anti-PD-L1 treatments can directly attenuate tumour-mediated exhaustion and effectively modulate the host anti-tumour immune response *in vivo*. In addition, compared with traditional therapy, PD-1/PD-L1 inhibitor monotherapy can significantly prolong survival without obvious side effects in the treatment of advanced NSCLC. Ideally, several biomarkers could be used to monitor the safety and effectiveness of anti-PD-1 and anti-PD-L1 treatments; however, the current lack of optimal prognostic markers remains a widespread limitation and challenge for further clinical applications, as does the possibility of immune-related adverse events and drug resistance. In this review, we aimed to summarise the latest progress in anti-PD-1/anti-PD-L1 treatment of advanced NSCLC, worldwide, including in China. An exploration of underlying biomarker identification and future challenges will be discussed in this article to facilitate translational studies in cancer immunotherapy.

Keywords: immunotherapy, non-small cell lung cancer (NSCLC), PD-1/PD-L1 inhibitors, predictive biomarkers, progress and challenges

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Introduction

Lung cancer is the leading cause of cancer-related deaths and is associated with a poor prognosis.^{1,2} Non-small-cell lung cancer (NSCLC) accounts for 85% of all lung cancers, 15–30% of which are squamous-cell carcinomas (SQCs) of the lung.^{3,4} The pathogenic genes of NSCLC have been discovered and widely investigated, and genotyping-based targeted therapy has been developed as one of the most successful therapeutic methods.^{5–8} At present, with a greater understanding of tumour immunity, targeted molecular therapy and immunotherapy have become important strategies for the conventional treatment of NSCLC.^{9–12} In recent years, the successful application of immune-checkpoint inhibitors (ICIs) of cytotoxic

T-lymphocyte antigen-4 (CTLA-4), and anti-programmed cell-death protein 1 (PD-1) and programmed cell-death protein ligand 1 (PD-L1) in various advanced cancers has attracted widespread attention in the field of immuno-oncology.^{13–18} Two antibodies against PD-1, nivolumab and pembrolizumab, and two antibodies against PD-L1, atezolizumab and durvalumab, have been approved by the European Medicines Agency (EMA) and the US Food and Drug Administration (FDA) for treating advanced NSCLC.^{19–24} The PD-1/PD-L1 inhibitors licensed for NSCLC treatment are listed in Table 1.

In this review, we aimed to outline the recent development of PD-1/PD-L1 inhibitors as first-,

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Table 1. The PD-1/PD-L1 inhibitors licensed for NSCLC treatment.

Checkpoint	Blocking agent	IgG isotype and characteristics	Clinical stage	Manufacturer
PD-1	Pembrolizumab	Humanized IgG4 mAb	EMA, FDA approved for first-line and second-line NSCLC treatment	Merck
	Nivolumab	Humanized IgG4 mAb	FDA approved for second-line NSCLC	Bristol-Myers Squibb
	Toripalimab	Humanized IgG4 mAb	Clinical trial ongoing	Shanghai Junshi
	Sintilimab	Fully human IgG4 mAb	Clinical trial ongoing	Innovent Biologics
	Camrelizumab	Humanized IgG4 mAb	Clinical trial ongoing	Hengrui Medicine
	Tislelizumab	Humanized IgG4 mAb	Clinical trial ongoing	BeiGene
	PDR001	Humanized IgG4 mAb	Clinical trial ongoing	Novartis
	REGN2810	Humanized IgG4 mAb	Clinical trial ongoing	Regeneron-Sanofi
	PD-L1	Atezolizumab	High-affinity human IgG1	FDA approved for second-line NSCLC
Durvalumab		Human IgG1 mAb	FDA approved for treatment of unresectable stage III NSCLC without relapse after chemoradiation	AstraZeneca
BMS-936559		Fully high-affinity human IgG4	Clinical trial ongoing	Bristol-Myers Squibb

EMA, European Medicines Agency; FDA, US Food and Drug Administration; IgG, immunoglobulin G; mAb, monoclonal antibody; NSCLC, non-small-cell lung cancer; PD-1, programmed cell-death protein 1; PD-L1, programmed cell-death ligand 1.

second-, and third-line treatments, and to discuss driver-gene mutation in NSCLC patients. Moreover, we summarised current and ongoing clinical trials of PD-1 and PD-L1 inhibitors in China. In the absence of optimal prognostic biomarkers, we elaborate on the presence of immune-related adverse events (irAEs), primary and acquired drug resistance, and the challenge of administering immunotherapy for the treatment of advanced NSCLC.

Regulatory mechanisms of PD-1/PD-L1 in NSCLC

Modulation of immune checkpoints is an important mechanism for evading immune surveillance by inhibiting activated T cells.²⁵ For example, PD-1⁺ immune cells can interact with PD-L1⁺ malignant tumour cells or other host cells. Mechanistically, immune escape might occur by interfering with related anti-tumour functions.²⁶ In addition, PD-1 is expressed by CD4⁺ T cells, CD8⁺ T cells and some other immune cell types. PD-1 interaction with its ligand, PD-L1, can weaken the effector functions of T lymphocytes such as activation, proliferation and survival,

promote antigen-specific T-cell apoptosis, and affect CD4⁺ and CD8⁺ T cells, natural killer (NK) cells, and CD11c⁺ M1 macrophage functions to promote tumour development.^{27–29} Antibodies that suppress the interaction between PD-1 and PD-L1 can largely improve and restore the functionality of exhausted T cells and enhance their anti-tumour immune response.³⁰ PD-1 is highly expressed by regulatory T cells (Tregs) and has a significant inhibitory effect on tumour immunity.³¹ The immune-related regulatory mechanism of action of PD-1/PD-L1 inhibitors is shown in Figure 1. Modulation of PD-1/PD-L interactions effectively inhibits the anti-tumour response by weakening the inhibitory activity of Tregs.³² In addition to T-cell immunity, tumour-associated macrophages,^{33,34} as well as NK cells and dendritic cells in the tumour microenvironment (TME) can enhance anti-tumour effects.^{35–37}

PD-1/PD-L1 inhibitor therapy in advanced NSCLC

Although there are many ways to treat NSCLC, such as radiotherapy, platinum-based chemotherapy, surgical resection and targeted molecular

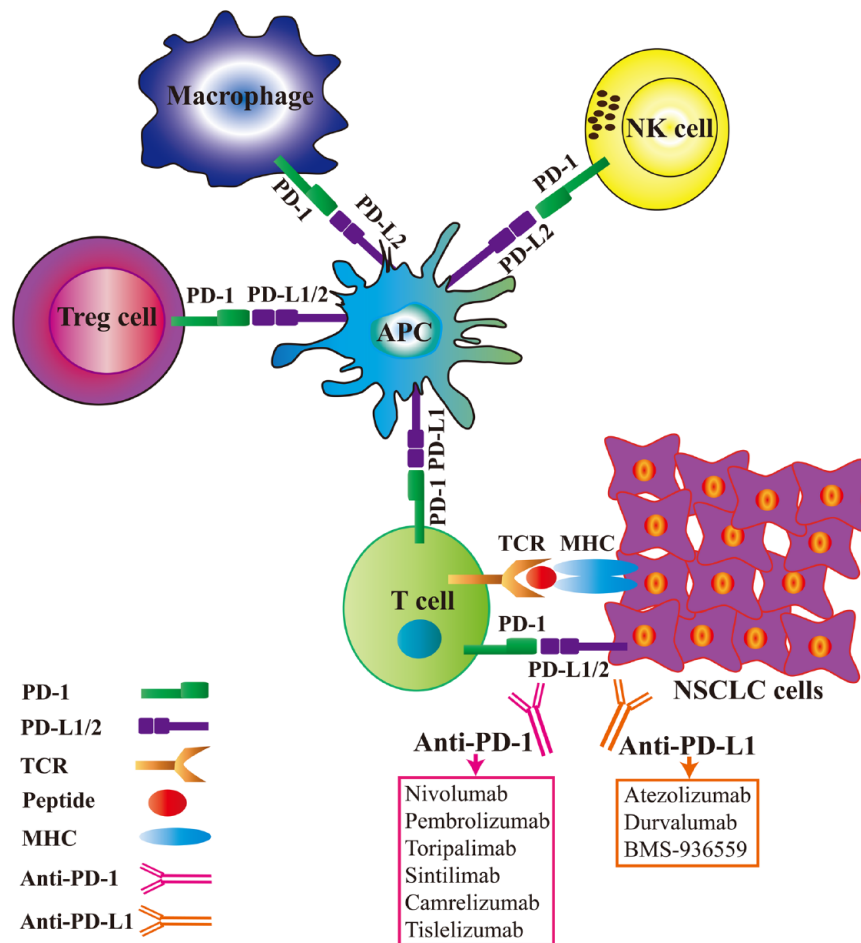


Figure 1. Regulatory mechanism of programmed cell-death protein 1 (PD-1)/programmed cell-death ligand 1 (PD-L1) inhibitors.

The PD-1 receptor is expressed on activated T cells, macrophages, regulatory T cells (Tregs) and natural killer (NK) cells. Binding of PD-1 to its ligands, PD-L1 or PD-L2, results in suppression of proliferation and inhibition of the immune response of T cells. Anti-PD-1 antibodies or PD-L1 inhibitors reverse this process, resulting in enhanced anti-tumour immune responses. APC, antigen-presenting cell; MHC, major histocompatibility complex; NSCLC, non-small-cell lung cancer; TCR, T cell receptor.

therapy, the long-term efficacy of these treatments is still not ideal, and the 5-year survival rate of patients with NSCLC is less than 18%.³⁸ Thus, there is an urgent demand for novel clinical therapies that are associated with less toxicity and higher efficacy to further prolong the survival time of patients. At present, some anti-PD-1/PD-L1 antibodies have been approved by the FDA for clinical treatment of advanced NSCLC;³⁹ these clinical studies will have a very important role in guiding long-term clinical practice.

PD-1/PD-L1 inhibitors as first-line treatment of advanced NSCLC

Multiple KEYNOTE trials (either alone or as part of a combined study) are being investigated

using PD-1/PD-L1 inhibitors in various tumour types, with a focus on NSCLC. In the open-label phase Ib KEYNOTE-001 trial, patients with advanced NSCLC expressing PD-L1 in tumour tissues who were randomly assigned to the pembrolizumab group exhibited an objective response rate (ORR) of 27%, a median progression-free survival (mPFS) of 6.2 months [95% confidence interval (CI) 4.1–8.6], and a median overall survival (mOS) of 22.1 months (95% CI 17.1–27.2). The ORR, 1-year mPFS, and 1-year mOS were significantly higher in patients with a PD-L1 tumour proportion score (TPS) $\geq 50\%$ than in the overall population. The results showed that pembrolizumab was well tolerated; only 12 cases (11.9%) experienced grade 3 treatment-related adverse events (AEs), and there

were no treatment-induced deaths.⁴⁰ The 3 years of data from KEYNOTE-001 showed that 79.7% patients with a PD-L1 TPS $\geq 50\%$ reached the mOS of 34.9 months (95% CI 20.3–not reached).⁴¹ The KEYNOTE-189 study is a phase III clinical trial of non-squamous NSCLC patients without first-line sensitised epidermal growth-factor receptor (EGFR) or anaplastic lymphoma kinase (ALK) mutations. These patients received platinum-based drugs plus either pembrolizumab or placebo. The mPFS of the monoclonal antibody (mAb) group was significantly longer than the control group (8.8 *versus* 4.9 months, respectively; $p < 0.001$), and the incidences of grade 3 or higher AEs in the mAb group and the control group were 67.2% and 65.8%, respectively.⁴² The update data from the KEYNOTE-189 study (23.1 months follow up) showed mOS was 22.0 months (95% CI 19.5–25.2) in the pembrolizumab–combination group *versus* 10.7 months (95% CI 8.7–13.6) in the placebo–combination group [hazard ratio (HR) 0.56; 95% CI 0.45–0.70]; the mPFS were 9.0 months (95% CI 8.1–9.9) and 4.9 months (95% CI 4.7–5.5), respectively (HR 0.48; 95% CI 0.40–0.58).⁴³

For patients with advanced squamous NSCLC, the KEYNOTE-407 trial indicated that concomitant administration of carboplatin plus paclitaxel and pembrolizumab provided longer mOS and mPFS than traditional chemotherapy alone.⁴⁴ The efficacy outcome from the protocol-specified final analysis of KEYNOTE-407 showed that, after a median follow up of 14.3 months, pembrolizumab plus chemotherapy exhibited a clinically meaningful improvement over placebo plus chemotherapy for mOS (17.1 *versus* 11.6 months) and mPFS (8.0 *versus* 5.1 months).⁴⁵

Another phase I/II trial, KEYNOTE-021, was designed to assess the efficacy and frequency of grade 3–4 AEs associated with combination treatment with platinum-based doublet chemotherapy with pembrolizumab. The results demonstrated that combination chemotherapy with pembrolizumab was associated with fewer grade 3–4 AEs compared with platinum-based chemotherapy or pembrolizumab monotherapy.⁴⁶ A randomised, open-label, phase II, multi-cohort study, KEYNOTE-021, was performed with 123 patients with primary stage IIIb, or IV NSCLC without EGFR or ALK mutations. The mPFS of the pembrolizumab plus chemotherapy group was 13.0 months (95% CI 8.3–not reached), whereas the mPFS of the group receiving chemotherapy

alone was only 8.9 months (95% CI 4.4–10.3).⁴⁷ After a median time of 49.4 months from KEYNOTE-021, ORR (58% *versus* 33%) and mPFS (24.5 *versus* 9.9 months; HR 0.54; 95% CI 0.35–0.83) was improved with the pembrolizumab combination compared with chemotherapy, regardless of PD-L1 status; the median OSs were, respectively, 34.5 *versus* 21.1 months (HR 0.71; 95% CI 0.45–1.12).⁴⁸

In the KEYNOTE-024 trial, compared with platinum-based chemotherapy alone, pembrolizumab significantly improved OS (30.0 *versus* 14.3 months), PFS (10.3 *versus* 6.0 months), and ORR (44.8% *versus* 27.8%) in the first-line therapy of advanced NSCLC cases with a PD-L1 TPS $\geq 50\%$.^{49,50} Moreover, pembrolizumab was associated with longer mOS and mPFS in contrast to chemotherapy alone in patients with a high PD-L1 TPS $\geq 50\%$ in the KEYNOTE-042 trial.⁵¹ A real-world study in Japan, HOPE-001, analysed clinical data from patients receiving pembrolizumab as first-line treatment for NSCLC. The ORR, mPFS, and mOS were 51.2%, 8.3 months (95% CI 6.0–10.7), and 17.8 months (95% CI 17.8–not reached), respectively,⁵² similar to the results of the KEYNOTE-024 and KEYNOTE-042 trials.

In the CheckMate 012 trial, 52 patients received nivolumab until disease progression or toxicity was experienced. The ORR was 28% in patients regardless of the tumour expression level of PD-L1, and the mPFS and mOS were 3.6 months (95% CI 0.1–28.0) and 19.4 months (95% CI 0.2–35.8), respectively.⁵³ In another CheckMate 012, open-label, phase I trial, 78 patients who received either nivolumab every 2 weeks plus ipilimumab every 12 weeks, or nivolumab every 2 weeks plus ipilimumab every 6 weeks, were assessed. The use of nivolumab plus ipilimumab as first-line treatment resulted in good tolerability and safety, and encouraging clinical efficacy, with a better ORR in each group.⁵⁴ Based on CheckMate 012, in the CheckMate 227 trial, nivolumab plus ipilimumab as first-line therapy significantly prolonged the mPFS compared with chemotherapy alone in patients with advanced NSCLC.^{55,56} CheckMate 568, a phase II trial of nivolumab plus ipilimumab in NSCLC patients, defined a cut-off of the tumour mutational burden (TMB) of at least 10 mutations per base pair for selecting patients most likely to respond to treatment, regardless of the PD-L1 expression level.⁵⁷ A TMB ≥ 10 was associated with a high

ORR and mPFS. However, in the CheckMate 026 study, compared with first-line chemotherapy alone in advanced NSCLC patients, patients that were treated with antibody therapy did not obtain significant survival benefits.⁵⁸

Atezolizumab and durvalumab are engineered and humanised mAbs against PD-L1 immunoglobulin G1 (IgG1), which can combine with PD-L1 to block the PD-1/PD-L1-mediated signalling pathway. BIRCH was designed to measure the efficacy of atezolizumab in advanced NSCLC as first-, second- and third-line therapies. Of 139 patients with advanced NSCLC who underwent first-line treatment with atezolizumab, the ORR, mPFS and mOS were 22%, 5.4 months (95% CI 3.0–6.9) and 20.1 months (95% CI 20.1–not reached), respectively.⁵⁹ IMpower150 was an open-label phase III study that aimed to assess the therapeutic efficacy of atezolizumab combined with bevacizumab and chemotherapy as first-line treatment in patients with advanced NSCLC. Patients received a combination therapy of atezolizumab + carboplatin + paclitaxel (ACP), bevacizumab + carboplatin + paclitaxel (BCP), or atezolizumab + BCP (ABCP) every 3 weeks for a total of four to six cycles. Following this, they then received atezolizumab, bevacizumab, or both, for maintenance treatment. Regardless of the expression levels and genetic status of PD-L1, the mPFS and mOS of patients with metastatic non-squamous NSCLC in the ABCP group were significantly improved when compared with the ACP or BCP group.⁶⁰ In the IMpower132 trial, atezolizumab combined with carboplatin and the chemotherapeutic agent pemetrexed significantly decreased mortality and improved the mPFS and mOS compared with chemotherapy alone in patients with advanced non-squamous NSCLC.⁶¹ The IMpower130 trial reported an obvious improvement in mOS and mPFS, and no change in AEs were found in the group receiving atezolizumab plus chemotherapy compared with the group receiving chemotherapy alone as the first-line treatment of patients with metastasised non-squamous NSCLC without driver gene mutations.⁶² The IMpower131 study showed that an atezolizumab plus platinum-based chemotherapy significantly improved mPFS in patients with first-line squamous NSCLC; however, the mOS was similar between groups.⁶³ Impower110 was performed to analyse the efficacy and safety of atezolizumab, as compared with platinum-based chemotherapy, as a first-line treatment for patients with metastatic NSCLC. The primary

measurement was the OS among patients in the intention-to-treat population. This study found that the mOS was longer in the atezolizumab group than in the chemotherapy group (20.2 *versus* 13.1 months; HR for death 0.59; 95% CI 0.40–0.89; $p=0.01$) as was mPFS (8.1 *versus* 5.0 months; HR for death 0.63; 95% CI 0.45–0.88) in the population with the highest expression of PD-L1. This study indicated that atezolizumab treatment results in significantly longer OS than platinum-based chemotherapy among patients with NSCLC with high PD-L1 expression, regardless of histologic type.⁶⁴ Recently, an ongoing APPLE study (WJOG11218L) was performed in Japan to evaluate the additional effect of bevacizumab administered with platinum combination therapy and atezolizumab in patients with advanced non-squamous NSCLC.⁶⁵

In a phase Ib trial, durvalumab and the anti-CTLA-4 antibody tremelimumab were administered to patients with advanced NSCLC. Regardless of the level of PD-L1 expression, the ORR was 23% in the group that received durvalumab combined with tremelimumab.⁶⁶ The MYSTIC trial was an open-label, phase III trial of first-line treatment with durvalumab with or without tremelimumab *versus* platinum-based chemotherapy alone in patients with advanced NSCLC. Compared with chemotherapy alone, the durvalumab group exhibited a clinically significant improvement in mOS when tumour cell PD-L1 expression was $\geq 25\%$.⁶⁷ The SAKK study was performed to evaluate the safety of first-line durvalumab in patients with advanced NSCLC and a performance status of 2. The data demonstrate that using first-line durvalumab in patients with PS2 and metastatic NSCLC with PD-L1 expression $\geq 25\%$ resulted in an unexpectedly high number of fatal early events due to rapid tumour progression.⁶⁸ All the clinical data described here are summarised in Table 2.

PD-1/PD-L1 inhibitors as second-line treatment in advanced NSCLC

The human anti-PD-1 mAb nivolumab was the first to be approved as second-line therapy for stage IV NSCLC.⁶⁹ CheckMate 017, a randomised, open-label, phase III study assessed the efficacy and safety of nivolumab for advanced squamous NSCLC; greater efficacy was obtained during or after first-line chemotherapy. Regardless of PD-L1 expression levels, the mOS, ORR, and mPFS were markedly better in the nivolumab

Table 2. Efficacy of anti-PD-1/PD-L1 agents for advanced NSCLC in first line.

Study	Histology	Phase	PD-L1 TPS (%)	Arms	Case number	ORR% (95% CI or p value)	mPFS (95% CI or p value, months)	mOS (95% CI or p value, months)	Grade 3-5 AE%	References
KEYNOTE-001	Squamous or non-squamous	Ib	≥50	Pembrolizumab	27	51.9 (31.9-71.3)	12.5 (6.2-NR)	NR (22.1-NR)	11.9	Hui <i>et al.</i> ⁴⁰
			1-49		52	17.3 (8.2-30.3)	4.2 (3.1-6.4)	19.5 (10.7-22.2)		
			<1		12	8.3 (0.2-38.5)	3.5 (2.1-19.0)	14.7 (3.4-NR)		
KEYNOTE-189	Non-squamous	III	Any	Pembrolizumab plus chemotherapy	410	47.6 (42.6-52.5)	8.8 (7.6-9.2)	NR	67.2	Gandhi <i>et al.</i> ⁴²
			Any	Chemotherapy	206	18.9 (13.8-25.0)	4.9 (4.7-5.5)	11.3 (8.7-15.1)	65.8	
KEYNOTE-189	Non-squamous	III	Any	Pembrolizumab plus chemotherapy	410	48.0 (43.1-53.0)	9.0 (8.1-9.9)	22.0 (19.5-25.2)	71.9	Gadgeel <i>et al.</i> ⁴³
			Any	Chemotherapy	206	19.4 (14.2-22.0)	4.9 (4.7-5.5)	10.7 (8.7-13.6)	66.8	
KEYNOTE-407	Squamous	III	Any	Pembrolizumab plus chemotherapy	278	57.9 (51.9-63.8)	6.4 (6.2-8.3)	15.9 (13.2-NR)	69.8	Paz-Ares <i>et al.</i> ⁴⁴
			Any	Chemotherapy	281	38.4 (32.7-44.4)	4.8 (4.3-5.7)	11.3 (9.5-14.8)	68.2	
KEYNOTE-407	Squamous	III	Any	Pembrolizumab plus chemotherapy	278	62.6 (56.6-68.3)	8.0 (6.3-8.4)	17.1 (14.4-19.9)	74.1	Paz-Ares <i>et al.</i> ⁴⁵
			Any	Chemotherapy	281	38.4 (32.7-44.4)	5.1 (4.3-6.0)	11.6 (10.1-13.7)	69.6	
KEYNOTE-021	Squamous or non-squamous	I/II	Any	Pembrolizumab plus carboplatin plus paclitaxel	25	48	10.3 (6.1-14.6)	21.4 (10.5-NR)	40	Gadgeel <i>et al.</i> ⁴⁶
			Any	Carboplatin plus paclitaxel plus bevacizumab	25	56	7.1 (4.2-14.3)	16.7 (8.5-NR)	42	
			Any	Carboplatin plus pemetrexed	24	75	10.2 (6.5-13.9)	16.7 (13.9-29.2)	46	
KEYNOTE-021	Non-squamous	II	Any	Pembrolizumab and chemotherapy	60	55 (42.0-68.0)	13.0 (8.3-NR)	NR	39	Langer <i>et al.</i> ⁴⁷
			Any	Chemotherapy alone	63	29 (18.0-41.0)	8.9 (4.4-10.3)	NR	26	

(Continued)

Table 2. (Continued)

Study	Histology	Phase	PD-L1 TPS (%)	Arms	Case number	ORR% (95% CI or p value)	mPFS (95% CI or p value, months)	mOS (95% CI or p value, months)	Grade 3–5 AE%	References
KEYNOTE-021	Non-squamous	II	Any	Pembrolizumab and chemotherapy	60	58	24.5 (9.7–36.3)	34.5 (24.0–NR)	39	Awad et al. ⁴⁸
				Chemotherapy alone	63	33	9.0 (6.2–15.2) HR: 0.54; [0.35–0.83]	21.1 (14.9–35.6) HR: 0.71; [0.45–1.12]	31	
KEYNOTE-024	Squamous or non-squamous	III	≥50	Pembrolizumab	154	44.8 (36.8–53.0)	10.3 (6.3–NR)	30.0 (18.3–NR)	31.2	Reck et al. ^{49,50}
				Chemotherapy	151	27.8 (20.8–35.7)	6.0 (4.2–6.2) p < 0.001	14.2 (9.8–19.0) p = 0.002	53.3	
KEYNOTE-042	Squamous or non-squamous	III	≥50	Pembrolizumab	299	39	7.1 (5.9–9.0)	20.0 (15.4–24.9)	18	Mok et al. ⁵¹
				Chemotherapy	300	32	6.4 (6.1–6.9) p = 0.0170	12.2 (10.4–14.2) p = 0.0003	41	
HOPE-001	Squamous or non-squamous	Real-world study	≥50	Pembrolizumab	213	51.2	8.3 (6.0–10.7)	17.8 (17.8–NR)	18.3	Tamiya et al. ⁵²
CheckMate 012	Squamous or non-squamous	I	Any	Nivolumab	52	23 (12.0–52.0)	3.6 (0.1–28.0)	19.4 (0.2–35.8)	19	Gettinger et al. ⁵³
CheckMate 012	Squamous or non-squamous	I	Any	2 weeks nivolumab plus 12 weeks ipilimumab	38	47 (31–64)	NA	NR	37	Hellmann et al. ⁵⁴
				2 weeks nivolumab plus 6 weeks ipilimumab	40	38 (23–55)	NA	NR	33	
CheckMate 568	Squamous or non-squamous	II	Any	Nivolumab (TMB ≥ 10)	48	44	7.1 (3.6–11.3)	NR	29	Ready et al. ⁵⁷
				Nivolumab (TMB < 10)	50	12	2.6 (1.4–5.4)	NR		
CheckMate 026	Squamous or non-squamous	III	≥5	Nivolumab	211	26 (20–33)	4.2 (3.0–5.6)	14.4 (11.7–17.4)	18	Carbone et al. ⁵⁸
				Chemotherapy	212	33 (27–40)	5.9 (5.4–6.9) p = 0.25	13.2 (10.7–17.1) HR: 1.01; [0.80–1.30]	51	

(Continued)

Table 2. (Continued)

Study	Histology	Phase	PD-L1 TPS (%)	Arms	Case number	ORR% (95% CI or p value)	mPFS (95% CI or p value, months)	mOS (95% CI or p value, months)	Grade 3-5 AE%	References
BIRCH	Squamous or non-squamous	NA	≥5	Atezolizumab	139	22	5.4 (3.0-6.9)	20.1 (20.1-NR)	40	Peters <i>et al.</i> ⁵⁹
IMpower150	Non-squamous	III	Any	ACP BCP ABCP	402 400 400	49.2 53.5 69.3	7.0 6.8 8.3 (p<0.001)	18.6 14.7 19.2 (p=0.02)	NA 47.7 57.7	Socinski <i>et al.</i> ⁶⁰
IMpower132	Non-squamous	III	Any	APP PP	292 286	49.6 32.2	7.6 (6.6-8.5) 5.2 (4.3-5.6) p<0.0001	18.1 (13.0-NR) 13.6 (11.4-15.5) p=0.0797	53.6 39.1	Papadimitrakopoulou <i>et al.</i> ⁶¹
IMpower130	Non-squamous	III	Any	Atezolizumab plus chemotherapy	451	49.2	7.0 (6.2-7.3)	18.6 (16.0-21.2)	81.0	Planchard <i>et al.</i> ⁶²
IMpower131	Squamous	III	Any	Atezolizumab + CnP CnP	343 340	49.7 41.0	6.3 5.6 (p=0.0001)	14.2 13.5 (p=0.16)	69.2 58.4	Jotte <i>et al.</i> ⁶³
Impower110	Squamous or non-squamous	III	≥50	Atezolizumab Chemotherapy	107 98	38.3 28.6	8.1 5.0 HR: 0.63; (0.45-0.88)	20.2 (16.5-NR) 13.1 (7.4-16.5) p=0.01	33.9 56.7	Herbst <i>et al.</i> ⁶⁴

ABCP, atezolizumab plus BCP; AE, adverse event; ACP, atezolizumab plus carboplatin plus paclitaxel; APP, atezolizumab plus pemetrexed plus carboplatin or cisplatin; BCP, bevacizumab plus carboplatin plus paclitaxel; CI, confidence interval; C, confidence interval; CnP, carboplatin + nab-paclitaxel; HR, hazard ratio; mOS, median overall survival; mPFS, median progression-free survival; NA, not applicable; NSCLC, non-small-cell lung cancer; NR, not reached; ORR, objective response rate; PD-1, programmed cell-death protein 1; PD-L1, programmed cell-death ligand 1; PP, pemetrexed plus carboplatin or cisplatin; TMB, tumour mutational burden; TPS, tumour proportion score.

group than in the docetaxel group.^{69,70} The CheckMate 057 trial evaluated the efficacy of nivolumab in patients with advanced non-squamous NSCLC who experienced obvious disease progression after first-line treatment. Compared with docetaxel, nivolumab treatment resulted in a remarkably longer mOS, with a more favourable safety profile.^{71,72} In both the CheckMate 017 and CheckMate 057 trials, the 2-year OS rates of patients with squamous NSCLC treated with nivolumab compared with docetaxel were about 23% and 8%, respectively. In those with non-squamous NSCLC, the 2-year OS rates of those treated with nivolumab were 29% and 16% in the CheckMate 017 and CheckMate 057 trials, respectively.⁷³ The CheckMate 078 trial compared the safety and efficacy of nivolumab with docetaxel in patients with squamous or non-squamous NSCLC.⁷⁴ The results were consistent with those of the CheckMate 017 and CheckMate 057 trials. Another Argentinian study was conducted to explore the efficacy and safety of nivolumab in previously treated patients with advanced NSCLC; the results showed that nivolumab was well tolerated, with promising prognostic outcomes in patients with NSCLC who had undergone previous treatment.⁷⁵ A multicentre retrospective observational study in Japan was performed to evaluate the efficacy of nivolumab in second-line treatment in NSCLC patients. The mOS was 14.6 months (95% CI 12.3–15.9); 1-year survival rate was 54.3%; mPFS was 2.1 months (95% CI 1.9–2.4); and ORR was 20.5%. Subgroup analysis showed that PS, EGFR mutation status, smoking status, and PD-L1 were associated with the effectiveness of nivolumab.⁷⁶

KEYNOTE-010, an open-label, randomised phase II/III trial, included pretreated NSCLC patients with an expression level of PD-L1 $\geq 1\%$ in tumour cells who had received treatment with pembrolizumab or docetaxel. This study showed that treatment with pembrolizumab significantly prolonged mOS in patients with previously treated, advanced NSCLC, especially in those with high PD-L1 expression levels.⁷⁷ Patients in the KEYNOTE-010 trial were followed up to assess long-term outcomes. At a median follow-up time point of 42.6 months (range 35.2–53.2 months), those treated with pembrolizumab experienced a significantly improved mOS compared with those treated with docetaxel. The 3-year OS rates were 34.5% *versus* 12.7% (for those with PD-L1 TPS $\geq 50\%$) and 22.9% *versus* 11.0% (for those with PD-L1 TPS $\geq 1\%$),

respectively.⁷⁸ The KEYNOTE-025 trial assessed the safety and efficacy of pembrolizumab in patients with NSCLC who had been previously treated in Japan.⁷⁹ The results showed that pembrolizumab was associated with a generally high ORR in those with a PD-L1 expression level $\geq 50\%$. The phase I/II KEYNOTE-021 study evaluated combination therapy with pembrolizumab plus ipilimumab in patients with previously treated advanced NSCLC. This study showed that among patients who received pembrolizumab (2 mg/kg) plus ipilimumab (1 mg/kg), ORR was 30%, mPFS was 4.1 months (95% CI 1.4–5.8), and mOS was 10.9 months (95% CI 6.1–23.7) but was associated with meaningful toxicity.⁸⁰

A randomised phase III clinical trial, OAK, also showed that regardless of the expression level of PD-L1, the survival rate following atezolizumab treatment was significantly improved compared with docetaxel. Based on the OAK findings, atezolizumab has been approved by the FDA and the EMA for further treatment of those previously treated for advanced NSCLC. For long-term survivors in the atezolizumab group, the ORR was 14% and the mOS was 13.8 months (95% CI 11.8–15.7).^{81,82} In the POPLAR study, an open-label, phase II trial, atezolizumab treatment significantly improved OS compared with docetaxel in patients with NSCLC who were previously treated.⁸³ The BIRCH study also proved the efficacy of atezolizumab in patients with advanced NSCLC as second-line therapy. The ORR, mPFS, and mOS were 19%, 2.8 months (95% CI 1.5–3.9), and 15.5 months (95% CI 12.3–not reached), respectively.⁵⁹ It is well known that approximately 10% of patients with NSCLC are complicated with comorbid interstitial pneumonia, with a poor prognosis. The Thoracic Oncology Research Group 1936/AMBITION study is an ongoing, multicentre, single-arm, phase II trial to assess the safety and efficacy of atezolizumab for pretreated advanced/recurrent patients with NSCLC complicated with idiopathic chronic fibrotic interstitial pneumonia. We are looking forward to the results of this clinical study.⁸⁴ Another multicentre, phase II, open-label, single-arm VinMetAtezo trial was designed to assess the safety and efficacy of metronomic oral vinorelbine in combination with atezolizumab in advanced NSCLC progressing after first-line platinum-based chemotherapy. The primary outcome was PFS at 4 months.⁸⁵ In France, Marine *et al.* conducted a study to evaluate the

cost effectiveness of atezolizumab *versus* docetaxel as a second-line treatment for stage IV NSCLC. Their results indicated that atezolizumab was more efficacious and more costly than docetaxel in treating advanced NSCLC.⁸⁶ All studies that have assessed PD-1/PD-L1 inhibitors as second-line therapies are summarised in Table 3.

PD-1/PD-L1 inhibitors as third-line or greater treatment in advanced NSCLC

The BIRCH study examined the clinical efficacy of atezolizumab as third-line or greater therapy in advanced NSCLC. The mOS, mPFS, and ORR were 13.2 months (95% CI 10.3–17.5), 2.8 months (95% CI 1.5–3.9), and 18%, respectively as third-line treatment, suggesting that atezolizumab has good clinical efficacy as third-line therapy for advanced NSCLC.⁵⁹

The ATLANTIC trial was an open-label, phase II, multicentre clinical trial conducted to assess the efficacy of durvalumab as third-line or greater therapy for advanced NSCLC. The ORR was 12.2% in EGFR⁺ and ALK⁺ patients who had expression levels of PD-L1 $\geq 25\%$ and received durvalumab, which was lower than that of patients who were EGFR⁻ and ALK⁻ with the same expression level of PD-L1 $\geq 25\%$ (16.4%) or those with expression of PD-L1 $\geq 90\%$ (30.9%). The mPFS rates in the three groups were 1.9 months, 3.3 months, and 2.4 months, respectively.⁸⁷

ARCTIC was a phase III, randomised, open-label study that evaluated the efficacy of durvalumab with or without tremelimumab *versus* the standard of care for advanced NSCLC as a greater than third-line therapy. For patients with expression levels of PD-L1 $\geq 25\%$, durvalumab resulted in significant improvements in mOS and mPFS compared with the standard of care. For patients with expression of PD-L1 $< 25\%$, durvalumab combined with tremelimumab resulted in obvious improvements in mOS compared with the standard of care.⁸⁸ A detailed summary of the clinical trials of third-line or greater treatment of advanced NSCLC is shown in Table 3.

PD-1/PD-L1 inhibitors in cases of advanced NSCLC with driver-gene mutations

Inhibition of the PD-1/PD-L1 axis has resulted in favourable responses in NSCLC patients with EGFR mutations; however, the efficacy was significantly lower in these patients compared with

those without EGFR mutations.^{89,90} Although high expression levels of PD-L1 in patients with advanced NSCLC may be associated with EGFR mutations,⁹¹ studies have shown that EGFR-tyrosine kinase inhibitors (TKIs) downregulate the messenger ribonucleic acid (mRNA) expression of PD-L1, suggesting that EGFR-TKIs may activate T-cell immunocompetence in the TME of NSCLC patients with EGFR-mutations.^{92–94} A phase II clinical trial showed that pembrolizumab was less effective in advanced NSCLC patients who were naïve to TKIs and were PD-L1⁺, with EGFR mutations, even those with a PD-L1 expression level $\geq 50\%$; this indicated that anti-PD-1/PD-L1 treatment was not the ideal first choice for patients with EGFR mutations.⁹⁵

The phase I/II KEYNOTE-021 trial analysed the feasibility of administering erlotinib or gefitinib combined with pembrolizumab as first-line treatment in patients with stage IIIb/IV EGFR-mutant NSCLC. Pembrolizumab plus gefitinib combination therapy had to be continued because of severe AEs, and another study showed that pembrolizumab plus erlotinib administration did not improve the ORR when compared with pembrolizumab plus gefitinib.⁹⁶ The CheckMate 370 trial evaluated the safety of first-line treatment with nivolumab combined with crizotinib in patients with ALK⁺ NSCLC. Again, the combination treatment had to be discontinued, and enrolment was closed because of observed grade ≥ 3 hepatic toxicities.⁹⁷

A subgroup analysis of advanced NSCLC patients with EGFR mutations and ALK rearrangements was performed to assess the use of anti-PD-1/PD-L1 therapy as second-line treatment in the CheckMate 057, KEYNOTE-010, POLAR and OAK studies. Their results revealed that EGFR mutations and ALK rearrangements in patients with advanced NSCLC were associated with beneficial outcomes and prolonged mOS from anti-PD-1/PD-L1 treatment compared with chemotherapy when administered as second-line treatment.^{71,72,77,78,81–83} In a study assessing the effects of nivolumab, the ORR was significantly lower in patients with EGFR-mutant non-squamous NSCLC than in patients without the mutation (8.8% *versus* 19.6%, $p=0.007$).⁹⁸ It is well known that the acquired EGFR-T790M mutation is a major reason for patients developing resistance to EGFR-TKIs. A clinical study showed that of 25 NSCLC patients who received nivolumab following previous TKI administration, 8 patients

Table 3. Efficacy of anti-PD-1/PD-L1 agents for advanced NSCLC in second line and third line.

Study	Histologies	Phase	PD-L1 TPS (%)	Arms	Case number	ORR (95% CI or p value)	mPFS (95% CI or p value, months)	mOS (95% CI or p value, months)	Grade 3-4 AE%	References
Checkmate 017 (second)	Squamous	III	Any	Nivolumab	135	20 (14-28)	3.5 (2.1-4.0)	9.2 (7.3-13.3)	7	Brahmer et al. ⁶⁹ ; Reck et al. ⁷⁰
Checkmate 057 (second)	Non-squamous	III	Any	Docetaxel	137	9 (5-15) <i>p</i> = 0.008	2.8 (2.1-3.5) <i>p</i> < 0.01	6.0 (5.1-7.3) <i>p</i> < 0.001	55	Borghaei et al. ⁷¹ ; Reck et al. ⁷²
				Nivolumab	292	19 (15-24)	2.3 (2.2-3.3)	12.2 (9.7-15.0)	10	
Checkmate 078 (second)	Squamous or non-squamous	III	Any	Docetaxel	290	12 (7-17) <i>p</i> = 0.02	4.2 (3.5-4.9) <i>p</i> = 0.39	9.4 (8.1-10.7) <i>p</i> = 0.002	54	Wu et al. ⁷⁴
				Nivolumab	338	16.6 (12.2-21.0)	2.8 (2.4-3.4)	12.0 (10.4-14.0)	10	
Keynote-010 (second)	Squamous or non-squamous	II/III	≥1%	Pembrolizumab 2 mg/kg	345	18	3.9 (3.1-4.1) <i>p</i> = 0.07	10.4 (9.4-11.9) <i>p</i> = 0.0008	13	Herbst et al. ^{77,78}
				Pembrolizumab 10 mg/kg	346	18	4.0 (2.7-4.3) <i>p</i> = 0.004	12.7 (10.0-17.3) <i>p</i> < 0.0001	16	
				Docetaxel	343	9 (<i>p</i> = 0.0005)	4.0 (3.1-4.2)	8.5 (7.5-9.8)	35	
				Pembrolizumab 2 mg/kg	139	30	5.0 (4.0-6.5) <i>p</i> = 0.0002	14.9 (10.4-NR) <i>p</i> = 0.0002	NA	
				Pembrolizumab 10 mg/kg	151	29	5.2 (4.1-8.1) <i>p</i> < 0.0001	17.3 (11.8-NR) <i>p</i> < 0.0001	NA	
Keynote-025 (second)	Squamous or non-squamous	Ib	≥1%	Docetaxel	152	8 (<i>p</i> < 0.0001)	4.1 (3.6-4.3)	8.2 (6.4-10.7)	NA	Nishio et al. ⁷⁹
				Pembrolizumab	38	22 (10-38)	3.9 (2.0-6.2)	19.2 (8.0-26.7)	29	
Keynote-021 (≥2)	NSCLC	I/II	Any	Pembrolizumab plus ipilimumab	44	30	4.1 (1.4-5.8)	10.9 (6.1-23.7)	29	Gubens et al. ⁸⁰
OAK (second)	Squamous or non-squamous	III	Any	Atezolizumab	425	14	2.8 (2.6-3.0)	13.8 (11.8-15.7)	15	Rittmeyer et al. ⁸¹ ; von Pawel et al. ⁸²

(Continued)

Table 3. (Continued)

Study	Histologies	Phase	PD-L1 TPS (%)	Arms	Case number	ORR (95% CI or p value)	mPFS (95% CI or p value, months)	mOS (95% CI or p value, months)	Grade 3-4 AE%	References
				Docetaxel	425	13	4.0 (3.3-4.2) p=0.049	9.6 (8.6-11.2) p=0.0003	43	
POPLAR (second)	Squamous or non-squamous	II	Any	Atezolizumab	144	17 (11.0-23.8)	2.7	12.6 (9.7-16.4)	11	Fehrenbacher <i>et al.</i> ⁸³
				Docetaxel	143	15 (9.3-21.4)	3.0 HR: 0.94; (0.72-1.23)	9.7 (8.6-12.0) p=0.04	39	
BIRCH (second)	Squamous or non-squamous	NA	≥5%	Atezolizumab	268	19	2.8 (1.5-3.9)	15.5 (12.3-NR)	NA	Peters <i>et al.</i> ⁵⁹
BIRCH (third)	Squamous or non-squamous	NA	≥5%	Atezolizumab	252	18	2.8 (1.5-3.9)	13.2 (10.3-17.5)	NA	Peters <i>et al.</i> ⁵⁹
ATLANTIC (third)	Squamous or non-squamous	II	≥25%	EGFR and ALK positivity (durvalumab)	74	12.2 (5.7-21.8)	1.9 (1.803.6)	13.3 (8.1-NR)	6	Garassino <i>et al.</i> ⁸⁷
				EGFR and ALK negativity (durvalumab)	146	16.4 (10.8-23.5)	3.3 (1.9-3.7)	10.9 (8.6-13.6)	9	
				≥90% PD-L1 expression (durvalumab)	68	30.9 (20.2-43.3)	2.4 (1.8-5.5)	NR (9.5-NR)	18	
ARCTIC (≥3)	Squamous or non-squamous	III	≥25	durvalumab	62	35.5	3.8 (1.9-5.6)	11.7 (8.2-17.4)	9.7	Planchard <i>et al.</i> ⁸⁸
				Standard of care	64	12.5	2.2 (1.9-3.7) HR: 0.71 (0.49-1.04)	6.8 (4.9-10.2) HR: 0.63 (0.42-0.93)	44.4	
			<25	Durvalumab + tremelimumab	174	14.9	3.5 (2.3-4.6)	11.5 (8.7-14.1)	22.0	
				Standard of care	118	6.8	3.5 (1.9-3.9) p=0.056	8.7 (6.5-11.7) p=0.109	36.4	

AE, adverse event; ALK, anaplastic lymphoma kinase; CI, confidence interval; EGFR, epidermal growth-factor receptor; HR, hazard ratio; mOS, median overall survival; mPFS, median progression-free survival; NA, not applicable; NSCLC, non-small-cell lung cancer; NR, not reached; ORR, objective response rate; PD-1, programmed cell-death protein 1; PD-L1, programmed cell-death ligand 1; TMB, tumour mutational burden; TPS, tumour proportion score.

exhibited the T790M mutation, rendering them resistant to EGFR-TKI treatment, whereas 17 patients did not. The mPFS times were 1.3 months (95% CI 0.1–1.8) and 2.1 months (95% CI 1.3–3.4) for the T790M-positive and T790M-negative patients, respectively ($p=0.099$), and the mOS was not achieved at the follow up at 7.3 months. Overall, this study indicated that patients resistant to EGFR-TKI with negative expression of T790M probably benefited from subsequent treatment with nivolumab.⁹⁹ Hence, some studies have explored combination therapy in advanced NSCLC patients who are resistant to TKIs.^{100,101} Gettinger *et al.* performed a study to assess the efficacy of nivolumab combined with erlotinib in advanced NSCLC patients with EGFR mutations who were treated with an EGFR-TKI but no chemotherapy.¹⁰² Another multicentre, open-label, phase Ib dose-escalation clinical trial was performed with patients previously treated with an ALK-TKI or chemotherapy who had advanced, ALK-rearranged NSCLC.¹⁰³ The patients received either nivolumab plus ceritinib (450 mg daily) or nivolumab plus ceritinib (300 mg daily). The results showed that the ORR and mPFS were improved in the nivolumab plus ceritinib (450 mg) group; however, this combination of drugs was also associated with increased toxicity. Based on safety considerations, an alternative regimen is being investigated in which ceritinib is administered as a monotherapy for two cycles before the initiation of combination treatment with nivolumab.

One of the studies intended to identify a biomarker for patients most likely to positively respond to immunotherapy. Tumour deoxyribonucleic acid (DNA) was derived from advanced NSCLC patients receiving nivolumab or pembrolizumab therapy as second- or third-line treatment. The most common mutated genes observed in these patients were tumour protein p53 (TP53; 49%), Kirsten rat sarcoma 2 viral oncogene homolog (KRAS; 43%), and v-erb-b2 erythroblastic leukaemia viral oncogene homolog 2 (ERBB2; 13%). This study confirmed that patients with KRAS mutations experienced a longer mPFS and mOS than patients without KRAS mutations. Furthermore, patients with ERBB-family mutations failed to respond to treatment with PD-1 antibodies.¹⁰⁴ The efficacy data of anti-PD-1/PD-L1 for advanced NSCLC with driver gene mutations is shown in Table 4.

Application of PD-1/PD-L1 inhibitors in advanced NSCLC cases in China

Toripalimab (also known as JS001 or TAB001) is a recombinant human anti-PD-1 mAb developed by Shanghai Junshi Biotechnology Co., Ltd. that selectively inhibits PD-1 and PD-L1/PD-L2 interactions to facilitate T-cell activation.¹⁰⁵ Yang *et al.* conducted a phase I dose escalation and expansion clinical study to evaluate the safety and tolerability of toripalimab in patients with advanced or recurrent malignancies in China.¹⁰⁶ Thirty-three patients participated in this study, seven of which had NSCLC. The results showed that toripalimab was well tolerated at a dose of 10 mg/kg every 2 weeks, and no dose-limiting toxicity was apparent; in addition, long-term anti-tumour activity was achieved in patients with alveolar soft-part sarcomas.¹⁰⁶

Sintilimab is a human IgG4 mAb coupled to PD-1; it was jointly developed by Innovent Biologics and Eli Lilly and Company. Sintilimab was approved for treating patients with classical Hodgkin's lymphoma who had relapsed or were refractory after at least two rounds of systemic chemotherapy in China.¹⁰⁷ A phase Ib study was performed to assess the safety and efficacy of sintilimab as a neoadjuvant therapy for NSCLC; the results showed that sintilimab was well tolerated in NSCLC patients, with an encouraging pathologic response rate.¹⁰⁸

Camrelizumab, a PD-1 inhibitor, developed by Jiangsu Hengrui Pharmaceutical Co., Ltd., has recently been approved for the treatment of relapsed or refractory classical Hodgkin's lymphoma in China.¹⁰⁹ A study evaluated the efficacy and safety of camrelizumab combined with microwave ablation (MWA) in patients with metastatic NSCLC. The results indicated that the ORR was 33.3%, with two patients achieving a complete response (CR) and five patients achieving a partial response (PR). The mPFS was 5.1 months; OS was not reached. The most common AE related to camrelizumab administration was reactive capillary hyperplasia of the skin¹¹⁰ that could be alleviated by substitution with apatinib.¹¹¹

The anti-PD-1 antibody, tislelizumab, has demonstrated clinical tolerability and safety in treating solid tumours, including NSCLC in Chinese patients.¹¹² Toripalimab, sintilimab, camrelizumab and tislelizumab are all currently undergoing clinical trials for the treatment of various solid

Table 4. Efficacy of anti-PD-1/PD-L1 agents for advanced NSCLC with driver gene mutations.

Study	Therapy line	Phase	Driver gene mutant	PD-L1 TPS (%)	Arms	Case number	ORR (95% CI or p value)	mPFS (95% CI or p value, months)	mOS (95% CI or p value, months)	Grade 3-4 AE%	References
NCT02879994	First line	II	EGFR	≥1%	Pembrolizumab	11	9	3.9	NR	NA	Lisberg <i>et al.</i> ⁹⁵
KEYNOTE-021 (NCT02039674)	First line	I/II	EGFR	Any	Pembrolizumab plus gefitinib	7	14.3	1.4 (0.2-13.0)	13.0 (0.2-NR)	71.5	Yang <i>et al.</i> ⁹⁶
Checkmate 370	First line	I/II	ALK	Any	Pembrolizumab plus erlotinib	12	41.7	19.5 (3.0-19.5)	NR	33.3	
Garassino <i>et al.</i> ⁹⁸	≥2	NA	WT-EGFR	Any	Nivolumab Plus Crizotinib	13	38	NA	NA	38	Spigel <i>et al.</i> ⁹⁷
					Nivolumab	1293	19.6 (17.4-21.7)	3.0 (2.8-31)	11.0 (10.0-12.0)	6	Garassino <i>et al.</i> ⁹⁸
					EGFR-mutant	102	8.8 (3.3-14.3) p=0.007	3.0 (2.7-3.3) p=0.004	8.3 (2.2-14.4) p=0.046	7	
Haratani <i>et al.</i> ⁹⁹	Second line	NA	T790M-negative	Any	Nivolumab	17	24	2.1 (1.3-3.4)	NR	NA	Haratani <i>et al.</i> ⁹⁹
					T790M-positive	8	13 p=1.000	1.3 (0.1-1.8) p=0.099	NR	NA	
Gettinger <i>et al.</i> ¹⁰²	Second line	I	EGFR	Any	Nivolumab Plus erlotinib	21	15	5.1 (2.3-12.2)	18.7 (7.3-NR)	24	Gettinger <i>et al.</i> ¹⁰²
Felip <i>et al.</i> ¹⁰³	Second line	Ib	ALK	Any	Ceritinib 450 mg plus nivolumab	8	50 (16-84)	6.4 (0.8-13.7)	NA	93	Felip <i>et al.</i> ¹⁰³
					Ceritinib 300 mg plus nivolumab	12	25 (6-57)	3.7 (1.8-NR)	NA	82	

ABCP, atezolizumab plus bevacizumab plus carboplatin plus paclitaxel; AE, adverse event; ALK, anaplastic lymphoma kinase; CI, confidence interval; EGFR, epidermal growth-factor receptor; mOS, median overall survival; mPFS, median progression-free survival; NA, not applicable; NSCLC, non-small-cell lung cancer; NR, not reached; ORR, objective response rate; PD-1, programmed cell-death protein 1; PD-L1, programmed cell-death ligand 1; TPS, tumour proportion score; WT, wild type.

tumours, including advanced NSCLC; these are summarised in Table 5.

Predictive biomarkers in anti-PD-1/PD-L1 immunotherapy

In order to predict the response to immunotherapy, the biomarkers initially used by the researchers were the expression of PD-1 receptor and PD-L1, and the prediction method was assessed only by immunohistochemistry.¹¹³ More recently, with the development with newer technology, there are many available biomarkers that could be used to further predict the efficacy and prognosis of immunotherapy.^{113–117} These include PD-L1 expression, the number of tumour-infiltrating lymphocytes (TILs), the TMB, mutations in certain driver genes, microsatellite instability (MSI)/defective mismatch repair (dMMR), peripheral blood-based biomarkers and interferon gamma (IFN- γ) levels (Figure 2).

Level of PD-L1 expression and the number of TILs

Currently, the level of PD-L1 is commonly used as a tumour biomarker to evaluate the efficacy of anti-PD-1/PD-L1 therapy in patients with NSCLC.^{34,118} The KEYNOTE-024 and KEYNOTE-042 trials^{49–51} suggested that pembrolizumab could significantly improve the survival rate in patients with high expression levels of PD-L1. The KEYNOTE-001 study also indicated that patients with a PD-L1 TPS $\geq 50\%$ would likely experience better outcomes, including improved ORR, mPFS, and mOS.⁴⁰ Therefore, pembrolizumab has been approved by the FDA for advanced NSCLC patients without driver gene mutations as a first-line treatment. In a single-arm study assessing early ICIs in selected patients with high expression levels of PD-L1, Ito *et al.* found that ORR and mPFS could be used as an endpoint to predict the survival of patients with high PD-L1 expression levels.¹¹⁹ Teramoto *et al.* analysed PD-L1 immunohistochemistry in tumour patients at stage I–IIIa NSCLC; they found that high PD-L1 expression is strongly related to early-stage NSCLC and, therefore, is a representative biomarker for predicting a good prognosis.¹²⁰ Recently, Ricci *et al.* investigated 150 advanced NSCLC patients, they suggested PD-L1 expression with a cut-off of 50% can be effectively evaluated by the cytological smears.¹²¹ Another study showed that PD-L1 expression, CD8⁺ T cell infiltration, the TMB, and the

human leukocyte antigen (HLA) class I were outstanding predictive biomarkers of the anticipated response to anti-PD-1 immunotherapy.¹²² The mPFS was significantly improved in patients with both a high level of PD-L1 expression and TMB or in those without a loss of HLA class I molecules.

The TME plays a vital role in the growth and progression of cancer cells and distant metastasis, and affects the efficacy of immunotherapy in clinical oncology.¹²³ TILs mainly include NK cells and cytotoxic T cells, both of which are routinely assessed in histopathology reports as predictive biomarkers.^{124,125} In hyperprogressive disease patients, a high number of CD2⁻ CD4 T lymphocytes during the first and second line of immunotherapy was associated with a lower mOS.¹²⁶ A retrospective study by Chen *et al.*, assessed tumour PD-L1 expression and T-cell infiltration in NSCLC patients with rare EGFR mutations and found that the frequency of a high PD-L1 expression level and CD8⁺ TIL infiltration was 36.7%, with a worse prognosis.¹²⁷

TMB

The TMB is representative of the number of tumour mutations; the higher the frequency of mutations in somatic exonic regions, the greater the number of corresponding tumour-related antigens that are expressed.^{128,129} For patients with advanced NSCLC, a high TMB detected by next-generation sequencing seems to predict the lasting clinical effects of PD-1 or PD-L1 blockades.¹³⁰ This results in the subsequent increase in activation of lymphocytes to maintain an immunological anti-tumour response. A previous study evaluated the predictive role of the TMB in 76 NSCLC patients treated with anti-PD-1/PD-L1 therapy. Patients with a high TMB exhibited significantly longer mPFS and mOS.¹³¹ In a Chinese study, the TMB was analysed in NSCLC patients receiving anti-PD-1/PD-L1 treatment; the mPFS was significantly longer at 10.6 months in the high TMB group compared with just 3.9 months in the low TMB group ($p=0.0007$), and the mOS was 21.0 months in the high TMB group and just 11.6 months in the low TMB group ($p=0.0126$).¹³² The circulating tumour DNA (ctDNA)-related TMB was also assessed in NSCLC patients who received anti-PD-1 or anti-PD-L1 therapy;¹³³ however, higher ctDNA-related TMBs were obviously related to shorter mOS and mPFS, which indicated that a higher ctDNA TMB

Table 5. Application of PD-1/PD-L1 inhibitors in the advanced NSCLC in China.

Drug	ClinicalTrials.gov identifier	Phase	Condition	Therapy line	Design	Patients (n)	Recruiting status	Locations
Toripalimab	NCT03513666	II	Advanced NSCLC with EGFR-mutation positive and T790M negative	Second line	Toripalimab combined with pemetrexed plus carboplatin	40	Not yet	Jiangsu Province Hospital
Toripalimab	NCT04316351	II	IIIB/IV NSCLC with T790M-positive mutations failed to osimertinib therapy	Second line	Toripalimab (JS001) combined with pemetrexed and anlotinib	60	Not yet	Guangzhou Institute of Respiratory Disease
Toripalimab	NCT03856411	III	Advanced NSCLC	First line	Toripalimab injection or placebo combined with standard chemotherapy	450	Recruiting	Cancer Hospital Chinese Academy of Medical Sciences
Toripalimab	NCT03924050	III	Advanced NSCLC with TKI-resistant EGFR mutated	Second line	Toripalimab or placebo combined with chemotherapy	350	Recruiting	Shanghai Pulmonary Hospital
Sintilimab	NCT03798743	II	Advanced NSCLC	Second line	Sintilimab combined with docetaxel	30	Recruiting	Hunan Province Tumor Hospital
Sintilimab	NCT03765775	II	Advanced NSCLC with first-generation EGFR-TKI resistance along with T790M negative	Second line	Anlotinib combined with sintilimab	20	Recruiting	First Hospital of Shijiazhuang
Sintilimab	NCT04252365	II	Advanced NSCLC	First line	Sintilimab and pembrolizumab	20	Not yet	Guangdong Association of Clinical Trials
Sintilimab	NCT03812549	I	Advanced NSCLC	First line	Sintilimab combination with radiation	29	Recruiting	West China Hospital, Sichuan University
Sintilimab	NCT04124731	II	Advanced NSCLC	First line	Sintilimab combined with anlotinib	98	Recruiting	Tianjin Medical University Cancer Institute and Hospital
Sintilimab	NCT03802240	III	EGFR-mutant advanced non-squamous NSCLC	Second line	Sintilimab ± IBI305 combined with pemetrexed and cisplatin	600	Recruiting	Shanghai Chest Hospital
Sintilimab	NCT04213170	II	Advanced NSCLC with brain metastases	NA	Sintilimab combined with bevacizumab	60	Recruiting	Sun Yat-sen University
Sintilimab	NCT03830411	II	Nonsquamous NSCLC with wild-type EGFR	Second line	Sintilimab compared with docetaxel or pemetrexed	76	Recruiting	China Three Gorges University
Camrelizumab	NCT04211090	II	Nonsquamous NSCLC with brain metastases and without driver gene mutations	First line	Camrelizumab combined with pemetrexed/carboplatin	64	Recruiting	Sun Yat-sen University
Camrelizumab	NCT04203485	III	PD-L1-positive advanced NSCLC	First line	Camrelizumab combined with apatinib mesylate or camrelizumab alone	762	Recruiting	Jiangsu HengRui Medicine Co., Ltd
Camrelizumab	NCT04303130	II	Advanced squamous NSCLC	First line	Camrelizumab combined with endostar	52	Recruiting	Beijing Cancer Hospital
Camrelizumab	NCT04167774	II	Advanced NSCLC	Second line	Camrelizumab combined with paclitaxel	62	Recruiting	Sun Yat-sen University
Tislelizumab	NCT03663205	III	Advanced non-squamous NSCLC	First line	Tislelizumab with chemotherapy versus chemotherapy	334	Not yet	Anhui Provincial Hospital
Tislelizumab	NCT03594747	III	Advanced squamous NSCLC	First line	Tislelizumab with chemotherapy versus chemotherapy	360	Not yet	Anhui Provincial Hospital

EGFR, epidermal growth-factor receptor; NSCLC, non-small-cell lung cancer; PD-1, programmed cell-death protein 1; PD-L1, programmed cell-death ligand 1; TKI, tyrosine kinase inhibitor.

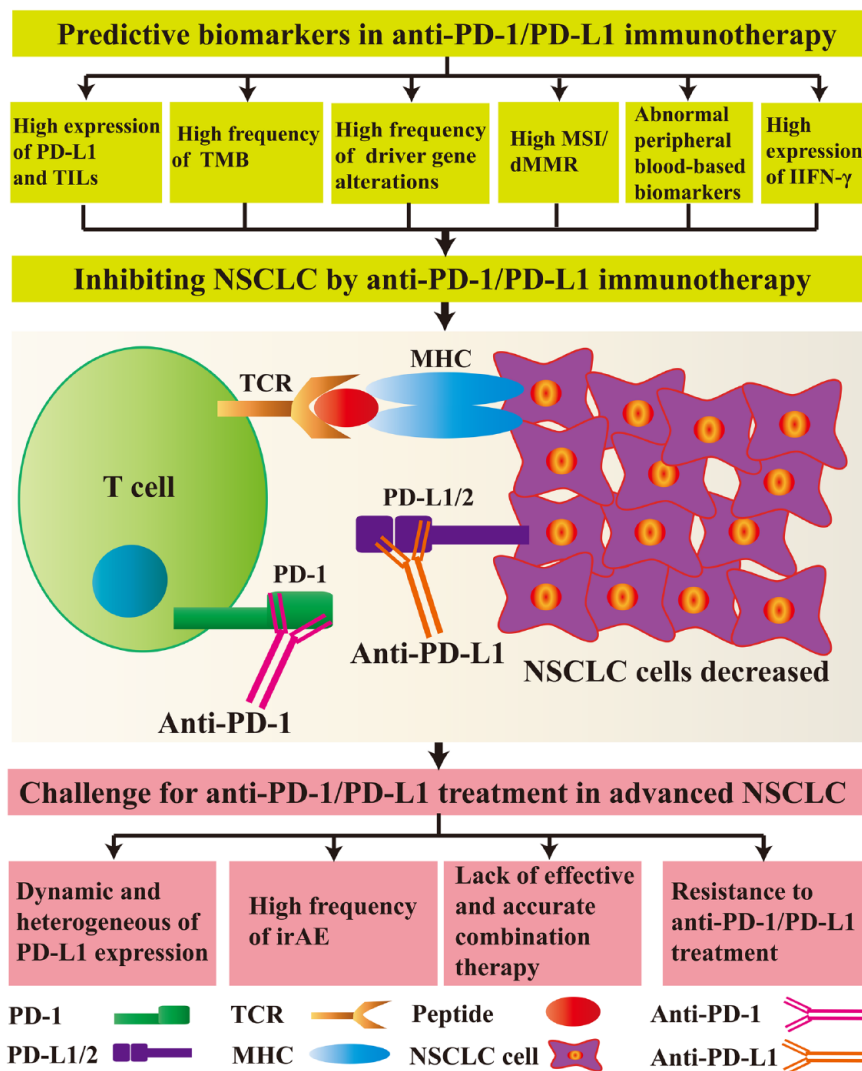


Figure 2. Predictive biomarkers and challenges facing anti-programmed cell-death protein 1 (PD-1)/programmed cell-death ligand 1 (PD-L1) treatment in advanced non-small-cell lung cancer (NSCLC). For inhibiting NSCLC cells, the predictive biomarkers include high expression levels of PD-L1, high numbers of tumour infiltrating lymphocytes (TILs), high tumour mutational burden (TMB), a high frequency of driver-gene alterations, high microsatellite instability (MSI)/defective mismatch repair (dMMR), abnormal peripheral blood-based biomarkers, and high expression levels of interferon gamma (IFN- γ). After administration of anti-PD-1 or anti-PD-L1 antibodies, the interaction between PD-1 and PD-L1/L2 is disrupted, inhibiting cell signalling pathways. T cells and other immune-related cells exert anti-tumour effects; however, there are still many unresolved problems in the clinical use of anti-PD-1/PD-L1 therapies, such as dynamic changes and heterogeneous differences in PD-L1 expression, a high frequency of immune-related adverse events (irAEs), the lack of effective and accurate combination therapies, and resistance to anti-PD-1/PD-L1 treatment. MHC, major histocompatibility complex; TCR, T cell receptor.

reflected worse clinical outcomes.¹³⁴ Another group detected TMB in plasma through blood-based cell-free DNA (cfDNA), and they were able to successfully identify NSCLC patients whose PFS was significantly improved after anti-PD-L1 therapy.¹³⁵ Chen *et al.* collected tumour tissue samples from 183 patients with NSCLC, and then assessed PD-L1 expression levels by immunohistochemical analysis, as well as TMB status to assess correlations between PD-L1

expression, TMB, and clinical pathological characteristics.¹³⁶ They found that in addition to quantifying existing PD-L1 expression levels, monitoring the TMB could be a valuable non-invasive biomarker for predicting the success of chemotherapy and more targeted treatments. This difference may be associated with the large amount of tumour heterogeneity in NSCLCs. A meta-analysis also found that the combination of CD8⁺ T-cell TILs, PD-L1 expression and TMB

was strongly related to a reliable prognosis for advanced NSCLC.¹³⁷ On the other hand, the precise cut-off values and methods of optimally monitoring TMBs need to be further explored in future clinical studies.

Driver-gene alterations

Next-generation sequencing technology has been able to identify thousands of mutations in large cohorts for various cancers.¹³⁸ Ono *et al.* investigated 242 patients with NSCLC to assess driver mutations using whole exome sequencing; they found that the existence of driver-gene alterations might be an independent factor that affects prognosis in NSCLC.¹³⁹ KRAS is one of the most common driver oncogenes associated with NSCLC.¹⁴⁰ A study investigated NSCLC-specific driver-gene alterations in the hope of identifying potential immunotherapy biomarkers. The most common driver genes were TP53 (49%), KRAS (43%) and ERBB2 (13%). Patients with KRAS mutations responded better to PD-1 inhibitors and exhibited improved mPFS and mOS than those without such mutations.¹⁰⁴ However, patients with ERBB-family mutations failed to respond to anti-PD-1 therapy. There was a case report of a 64-year-old non-smoking man with squamous NSCLC with co-mutations of TP53 and KRAS who received pembrolizumab combined with gemcitabine treatment; this patient achieved PR, which had persisted for over 7 months at the time of publication.¹⁴¹ However, another study retrospectively identified patients with NSCLC who were harbouring KRAS mutations who were treated with immunotherapy, and there was no significant difference between patients with or without KRAS mutations in terms of mPFS.¹⁴² Another study showed that the reason patients with KRAS mutations responded better to PD-1/PD-L1 inhibitors was related to tumour immunogenicity. NSCLC patients with KRAS mutations show good clinical outcomes following treatment with PD-1/PD-L1 inhibitors.¹⁴³

MSI/dMMR

In addition to PD-1/PD-L1 expression analysis, the presence of MSI is also a key reaction marker.¹⁴⁴ Studies have shown that genetic mutations in the dMMR pathway can lead to MSI, and high MSI is closely related to the increased immunogenicity and immune response produced by tumours. MSI is an excellent predictive

biomarker. Therefore, the FDA has approved pembrolizumab to treat unresectable solid tumours with high MSI or dMMR.¹⁴⁵ The DNA MMR pathway includes the mutL homolog 1 (MLH1), MSH (mutS homolog)-2, postmeiotic segregation increased 2 (PMS2) and MSH-6 proteins. When this pathway is defective (dMMR), it can be used as a means to select patients for immunotherapy. In addition, MSI frequency can also be used to predict NSCLC.¹¹⁷ Similarly, NSCLC patients with DNA polymerase epsilon (POLE) and DNA polymerase delta 1 (POLD1) gene mutations also have an objective clinical response to pembrolizumab treatment and a corresponding degree of association.^{146,147} Therefore, clinicians can further determine whether a patient can use the epidemiology of the patient's MSI/dMMR to predict NSCLC prognosis.¹⁴⁸

Peripheral blood-based biomarkers

Because biopsies of lung cancer tissue samples are difficult to obtain, some alternative methods include minimally invasive biopsy of peripheral blood for practical reasons.¹⁴⁹ Peripheral blood contains DNA, RNA, and proteins released from tumour tissues reflecting the dynamic changes in the TME. The level of ctDNA in the peripheral blood in the early stage of treatment provides valuable clues to the therapeutic effect. In various cancer types, including NSCLC, the detection of highly mutated ctDNA is positively correlated with a better clinical response to immune-checkpoint blockade.¹⁵⁰ Zhang *et al.* analysed individual peripheral T-cell clones of 110 peripheral T cells of NSCLC patients before and after the initiation of PD-1 blockade.¹⁵¹ They found that 25 genes were significantly changed during tumour progression. These analyses help to understand the dynamic changes of T-cell clones in the peripheral blood during PD-1 blockade of NSCLC. Tanizaki *et al.* investigated 134 patients with advanced or recurrent NSCLC treated with nivolumab to evaluate the association between survival and peripheral blood parameters; they found that a low absolute neutrophil count, high absolute lymphocyte count, and high absolute eosinophil count were closely related to a better clinical outcome in patients with NSCLC treated with nivolumab.¹⁵² In addition, Jiang *et al.* investigated 76 NSCLC patients to evaluate the relationship between survival and peripheral blood markers with ICI monotherapy; they found that a high platelet-to-lymphocyte ratio and low albumin in peripheral blood might be important indicators in

advanced NSCLC patients treated with nivolumab and durvalumab monotherapy.¹⁵³

IFN- γ

IFN- γ is a vital immune-regulatory molecule in the TME released by tumour-reactive T cells.¹⁵⁴ Recent studies have suggested that IFN- γ is an important driver of PD-L1 expression in tumours, and it can significantly improve the response to anti-PD-1 therapy.¹⁵⁵ The increased IFN- γ levels at baseline and at 3 months after anti-PD-1 therapy were significantly correlated with improved mOS.¹⁵⁶ Recent studies have shown that in 17 NSCLC patients treated with nivolumab, mPFS was obviously prolonged in those with high expression levels of IFN- γ , in contrast to those with low expression of IFN- γ (5.1 *versus* 2 months, respectively; $p=0.0124$). This indicated that IFN- γ is a vital biomarker for NSCLC patients treated with anti-PD-1 therapy.¹⁵⁷

Other potential biomarkers

Other potential biomarkers being investigated in anti-PD-1/PD-L1 immunotherapy on NSCLC include the pretreatment haemoglobin level,¹⁵⁸ serum vascular endothelial growth factor (VEGF) concentration,¹⁵⁹ serum lactate dehydrogenase level,¹⁶⁰ gut microbiome composition¹¹⁶ and neoantigen load.¹⁶¹

Challenges for anti-PD-1/PD-L1 treatment in advanced NSCLC

Currently, there are still many unresolved problems associated with the clinical use of anti-PD-1/PD-L1 therapies (Figure 2). First, PD-L1 expression is a predictive biomarker of outcomes of anti-PD-1/PD-L1 treatment, as it is correlated with the efficacy of PD-1/PD-L1 inhibitor therapy in NSCLC; however, this association can be quite variable. For example, the improvement in mOS by nivolumab and durvalumab treatment for NSCLC was independent of PD-L1 expression levels.^{71,162} In immunohistochemical analysis, the expression levels of PD-L1 are dynamic and heterogeneous, and this is further complicated by issues related to antibody specificity and detection techniques.¹⁶³ In the future, more valuable biomarkers may be identified by neoantigen prediction techniques, microbiota signatures, or high microsatellite instability (MSI), all of which should be further explored in clinical medicine.

The second challenge is the risk of irAEs associated with the use of anti-PD-1/PD-L1 monotherapy for advanced NSCLC. Although the safety profile of individual antibodies is favourable compared with standard chemotherapy; the frequency of irAEs in combination therapies was much higher (Tables 2–4). For example, 53.6% of patients who received atezolizumab plus chemotherapy developed irAEs, whereas only 29.1% of patients in the group receiving chemotherapy alone did.⁶¹ Therefore, better strategies to reduce the risks of serious irAEs need to be explored in future studies.

The third challenge of immunotherapy is deciding how to effectively and accurately select the best combination therapy for patients with advanced NSCLC. Anti-PD-1/PD-L1 treatment combined with chemotherapy is proven to have great potential for improving ORRs.^{164,165} Apart from chemotherapy, treatments with VEGF, CTLA-4, chimeric antigen receptor T cell, CRISPR-Cas9 gene editing or lymphocyte-activation-protein-3 therapies could provide additional choices for combination strategies in future clinical studies.^{17,166–169} Therefore, it is important to establish a standard to guide and determine the most appropriate combination therapy for treating advanced NSCLC.

The fourth challenge is that some patients who are resistant to anti-PD-1/PD-L1 therapy experience a poor prognosis after treatment. The underlying resistance mechanism is a very complicated process, and little is known about the potential tumour-related antigens associated with TMB resistance to anti-PD-1/PD-L1 therapy or the genetic determinants of a good prognosis.

Conclusion

To date, four anti-PD-1/PD-L1 drugs, including pembrolizumab, nivolumab, atezolizumab and durvalumab, have been registered and approved for the neoadjuvant, first-line and second-line treatment of NSCLC. With the approval of ICIs, immune-checkpoint therapy based on modulating PD-1/PD-L1 signalling provides more choices for treating advanced NSCLC patients. However, the ORR after ICI treatment is only 14–20% in NSCLC patients who are not identified beforehand as being likely to experience a good prognosis. Therefore, identification of predictive biomarkers is crucial for selecting patients with NSCLC who would most benefit from

anti-PD-1/PD-L1 therapy. Although the clinical application of anti-PD-1/PD-L1 therapy is still at an early stage, there are an increasing number of ongoing clinical trials to assess the efficacy of anti-PD-1/PD-L1 treatment. EGFR mutation rates and microbiome differences may affect the efficacy of anti-PD-1/PD-L1 treatment. ALK rearrangements in patients with advanced NSCLC are associated with beneficial outcomes and prolonged mOS from anti-PD-1/PD-L1 treatment compared with chemotherapy, when administered as second-line treatment. Patients with KRAS mutations experienced a longer mPFS and mOS than patients without a KRAS mutation using anti-PD-1/PD-L1 treatment, thus immunotherapy may be a promising strategy with KRAS mutation of NSCLC owing to regulating immune response. Therefore, with the continuous development of clinical trials, more retrospective and prospective clinical studies need to comprehensively evaluate more underlying biomarkers to improve treatment efficacy and enable more NSCLC patients to benefit from ICIs for anti-PD-1/PD-L1 therapy.

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Authors' contributions

JZ and JQ wrote the manuscript; QM, LC, LL, TC and PW collected the references and modified the manuscript; LC and JZ designed the manuscript and approved the final manuscript for publication.

Availability of data and materials

Please contact the corresponding author for data requests.

Conflict of interest statement

The authors declare that there is no conflict of interest.

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