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Immune Checkpoint Inhibitor Therapy in Cancer: Success versus Limitations

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ABSTRACT

The immune system possesses the capability to identify tumor cells and eradicate early malignant tumor cells. Thus, activating the immune system of cancer patients provides great therapeutic benefits. Inhibitory T-cell immune checkpoints play a vital role in tumor immune escape. Thus, immune checkpoint inhibitors (ICIs) have attracted attention in cancer immunotherapy. In ICI therapy, the therapeutic targets are the expressed immune checkpoints of T cells. Immune checkpoints induce T-cell dysfunction in cancer. However, ICIs or immunomodulators restore the antitumor actions of cytotoxic T cells by blocking immune checkpoints. ICIs have become desirable treatment options because of their broad range of activities and response rates ranging from 15% to 90% in several cancer types. Generally, ICIs also have favorable toxicity profiles. This paper will first delve deeper into the best-known immune checkpoints and then review ICIs that are attractive treatment options in immunotherapy.

Key words: Immune Checkpoint Inhibitor, T-cell, Immunotherapy, Cancer, Tumor

INTRODUCTION

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Cancer is considered the second leading cause of death of human beings worldwide after cardiovascular disease, and despite tremendous scientific developments, the cure for cancer is still challenging¹. The reasons behind this include lifestyle, method of eating, societal change, industrialization, environmental pollution, and inadequate treatment options ^{1,2}. Among the different etiological factors contributing to the genesis of cancer, societal modernization is regarded as a leading factor ^{1–4}. It is estimated that the number of deaths from cancer may increase to 11 million by 2030^{1,5}.

Treatment options such as surgery are effective for localized cancers only. Radiotherapy can play a vital role in cancer treatment if the cancer is not disseminated⁵. Chemotherapy is considered one of the primary treatment options for cancer. To date, various chemotherapeutic agents have been developed and are being used either alone or in combination to treat cancer^{5,6}. Also, the discovery of the chemotherapeutic agent, cisplatin, and the development of its various analogs have created optimism in chemotherapy ^{1,4}. However, the side effects of these chemotherapeutic agents restrict their successful use ^{1,3,5}. Again, nanoparticles, such as ZnO, have gained increased attention as chemotherapeutic agents because of their targeted action and minimal side effects⁵. Despite significant advances in understanding cancer etiology, ideal anticancer drugs or strategies are still missing ^{1,3}.

Established tumors require the employment of diverse mechanisms to suppress the antitumor response. The mechanisms include the upregulation of coinhibitory molecules called immune checkpoints, engaging immunosuppressive immune cells and inhibitory cytokine production⁷. Nevertheless, the immune system possesses the capability to identify tumor cells and eradicate early malignant tumor cells⁸. Thus, activating the immune system of cancer patients for therapeutic benefit has long been pursued by scientists⁹. Research on inhibitory T-cell immune checkpoints has realized this goal¹⁰. Furthermore, recent studies have shown that inhibitory T-cell immune checkpoints play a vital role in tumor immune escape. Thus, immune checkpoint inhibitors (ICIs) have attracted attention in cancer immunotherapy¹¹. As a result, the past decade has observed a quick transition in cancer treatment with the advent of immunother apy^{12} .

Immunotherapy is one oncologic treatment type undertaken to enhance host immunity to fight cancer¹³. It harnesses the immune system's capability to identify nonself-tumor antigens and constantly adapt to and detect new antigens¹⁴. In addition, it targets the development of tumor rejection capabilities, breaking the tumor-induced immune tolerance¹⁵. The advan-

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tages of immunotherapy include sustained surveillance, a low toxicity profile, and the ability to detect a small number of cancer cells¹⁶. Thus, immunotherapy is regarded as an alternative method of cancer treatment. Furthermore, the success of ICIs suggests that active immunotherapy can contribute to obtaining a longer-lasting response in cancer patients^{10,11,17}. In this paper, ICIs and their role in immunotherapy are discussed in depth with their prospects in medical science.

Literature research was performed in PubMed and Google Scholar with the following search terms: "immune checkpoint inhibitor" and "cancer" in combination with "melanoma," "metastasis," "ICI," "PD-1," "CTLA-4," "LAG-3," "TIM-3," "BTLA," "PD-L1," "PD-L2," and "toxicology."

IMMUNE CHECKPOINTS

Tumor-infiltrating lymphocytes (TILs), a kind of naturally occurring T cells, can identify tumor antigens^{13,18}. However, TILs, especially CD8⁺ T cells, cannot successfully eliminate cancerous cells when the effector function of infiltrating T cells is curtailed by immunosuppressive mechanisms of the tumor microenvironment¹⁹. Among these mechanisms, the upregulation of immune checkpoints has emerged as a major marker for the dysfunction of T cells in cancer²⁰. Thus, in ICI therapy, the therapeutic targets are the expressed immune checkpoints of the T cells. Immune checkpoints are coinhibitory or costimulatory molecules that induce T-cell dysfunction in cancer. However, ICIs or immunomodulators restore the antitumor actions of cytotoxic T cells by blocking immune checkpoints²¹. ICIs have become very appealing treatment options because of their broad range of activities and response rates ranging from 15% to 90% in several cancer types. Generally, ICIs also have favorable toxicity profiles²².

The best-known immune checkpoints corresponding to T-cell dysfunction along with controlling immune responses include PD-1 (Programmed Death 1), CTLA-4 (Cytotoxic T-lymphocyte-associated Antigen 4), LAG-3 (Lymphocyte-activation Gene 3), TIM-3 (T-cell Immunoglobulin and Mucin Domain 3), and BTLA (B- and T-lymphocyte Attenuator)^{20,21,23}. These best-known immune checkpoints will be discussed in depth in the following sections.

PD-1

PD-1 is a significant controller of tumor T-cell effector functions²⁴. This CD28 homolog is expressed in activated T cells, natural killer (NK) T cells, dendritic

cells (DCs), and activated B cells, to name a few. Furthermore, expression is induced by interleukin (IL)-21, IL-15, IL-7, and IL-2 on T cells 16,21,25-28. It was isolated from a T-cell hybridoma that was experiencing T-cell receptor (TCR) activation-induced cell death. Thus, it was called PD-19,29. This type I transmembrane protein comprises an intracellular domain with an ITIM (immunoreceptor tyrosine-based inhibitory motif) along with an ITSM (immunoreceptor tyrosine-based switch motif), a transmembrane domain, an immunoglobulin (Ig) superfamily domain, and a stalk of ~20 amino acids (aa) 11,30. The tyrosine of ITSM is necessary for PD-1 functioning in B and T cells²⁸. When ITIM and ITSM become phosphorylated, they recruit Src homology 2 (SH2)-containing tyrosine phosphatase 1 (SHP-1) along with SH2containing tyrosine phosphatase 2 (SHP-2). SHP-1 and SHP-2 reduce TCR signaling by dephosphorylating the CD3zeta chain^{21,30}.

On mouse chromosome 1 and human chromosome 2, PD-1 is produced by the *Pdcd1* gene. In both humans and mice, *Pdcd1* is composed of 5 exons. A short signal sequence and an Ig domain are produced by exons 1 and 2, respectively. Exon 3 is composed of the transmembrane domain and the stalk. A short, 12 amino acid sequences are encoded by exon 4, indicating the cytoplasmic domain's outset. Exon 5 contains long 3' UTR and C terminal intracellular residues³⁰. The biological significance of PD-1 pervades the immune response, such as tumor immunity, autoimmunity, transplantation immunity, infectious immunity, and allergy³¹.

PD-L1 is, in fact, a PD-1 ligand ^{11,30}. This type 1 transmembrane protein is also called CD274 and B7-H1⁸. In humans, it is expressed by, among others, macrophages, vascular endothelial cells, astrocytes, pancreatic islet cells, T cells, DCs, and B cells³². On mouse chromosome 19, PD-L1 is produced by the *Cd274* gene. However, in human chromosome 9, it is produced by the *Pdcd1* gene. *CD274* consists of 7 exons. Exon 1 possesses a 5' UTR, and exon 2 possess a signal sequence. Again, exons 3 and 4 possess an IgV-like and an IgC-like domain, respectively. Furthermore, exons 5 and 6 possess a transmembrane and intracellular domain, respectively. Exon 7 contains a 3' UTR and ~30 aa long intracellular residues³⁰.

PD-L2 is also a PD-1 ligand ³⁰. This type 1 transmembrane protein is also termed B7-DC or CD273^{8,30}. Its expression is limited to DCs, macrophages, and B cells and is encoded by the gene *Pdcd1lg2* neighboring the gene *Cd274*³². In mice, the *Pdcd1lg2* gene consists of 6 exons, while it consists of 7 in humans. Exon 1 is noncoding, and exon 2 possesses a signal sequence.

Furthermore, the IgV-like and IgC-like domains are produced by exons 3 and 4, respectively. Exon 5 consists of a transmembrane region, a short stalk, and the outset of the cytoplasmic domain. Exon 5 in mice possesses a stop codon that produces a 4 aa long cytoplasmic domain. Finally, the additional coding region in humans (exon 6 and 7) results in a 30 aa long cytoplasmic domain³⁰.

CTLA-4

CTLA-4, or CD152, is a significant controller of T-cell activation^{21,24}. This homolog of TCR CD28 is expressed on activated B cells in addition to activated CD4⁺ and CD8⁺ T-cell surfaces^{21,33}. It binds to ligands CD80 (also termed B7-1) and CD86 (or B7-2) with increased affinity compared to CD28^{8,33-35}. CTLA-4 binding with ligands with higher affinity than CD28 decreases CD28-dependent costimulation. Again, CTLA-4 can mediate inhibitory effects on the major histocompatibility complex (MHC)-TCR pathway directly 21,36. CTLA-4 can recruit SHP-2 along with PP2A to the intracellular YVKM domain. SHP-2 attenuates TCR signaling by dephosphorylating the CD3zeta chain. PP2A impairs TCR signaling by inhibiting downstream Akt phosphorylation^{21,24,32}. Actually, CTLA-4 was isolated from a mouse T-cell cDNA library. Hence, it was named CTLA-4, comprised of 3 introns and 4 exons, and encodes a type I transmembrane protein³⁴. CTLA-4 also shares a conserved motif of the MYPPPY amino acid sequence with CD28. This conserved motif is thought to be critical for ligand binding 34,36,37.

LAG-3

LAG-3, another type I transmembrane protein, is associated with the Ig superfamily and possesses 4 Iglike domains³⁸. This CD4-related inhibitory receptor is also known as CD223. It has a higher binding affinity for MHC II than for CD4, and is expressed in NK cells, activated CD4⁺ and CD8⁺ T cells, B cells, TILs, and T-regulatory cells (Tregs). On tolerant TILs, LAG-3 is coexpressed with PD-1. It further suppresses the activation of antigen-presenting cell (APC) binding with MHC II molecules 8,16,24,39. Again, it inhibits CD4⁺activation and decreases CD8⁺ cytotoxic function²⁰. This inhibitory function is dependent on signaling through the cytoplasmic domain motif KIEELE. In addition, LAG-3 transfection into T cells can provide a regulatory function. Thus, it is essential for the maximal function of Tregs⁸. However, the exact molecular mechanism of downstream signaling of LAG-3 is not entirely known³⁸.

TIM-3

TIM-3, another type I transmembrane protein, is expressed on monocytes, DCs, T cells, Tregs, and macrophages^{20,39,40}. Furthermore, it is found in activated human NK cells and suppresses the cytotoxicity of NK cells⁴¹. TIM-3 is associated with carcinogenesis, as its expression correlates with reduced survival, tumor invasion, and metastasis. It can be coexpressed with PD-1 on APCs²⁰. Several ligands, such as phosphatidylserine, CEACAM-1, galectin-9, and HMGB1, have been reported for TIM-3⁸. Also, TIM-3 causes CD4⁺ and CD8⁺ T cell apoptosis upon galectin-9 binding through the calcium-calpain-caspase-1 pathway²⁰. Together with PD-1 and other inhibitory receptors, it can mediate the exhaustion of CD8⁺T cells⁴¹.

BTLA

BTLA, or CD272, a glycoprotein of the Ig superfamily, has been identified as an inhibitory receptor^{20,42-44}. This type I transmembrane cosignaling receptor is structurally connected with CTLA-4 and PD-1^{40,45}. It is expressed on macrophages, DCs, T cells, B cells, and NK cells 20,46. In its cytoplasmic domain, it contains ITIMs and recruits both SHP-1 and SHP-2 upon phosphorylation. In addition, it negatively regulates both TCR and B-cell receptor signaling in vitro 42,44,46. For example, the herpes virus entry mediator (HVEM) transmits signals to BTLAexpressing cells and functions as a ligand⁴². HVEM has 4 cysteine-rich domains in its extracellular region that mediate the binding of HVEM with BTLA. Upon binding, HVEM transmits signals by TNF receptorassociated factor 2 to induce STAT3 phosphorylation, which results in the activation of NF-kB and other prosurvival signals⁴⁶. Again, the binding of BTLA with ligands decreases cytokine production and T-cell proliferation⁴⁰. Thus, BTLA is considered a possible target in immunotherapy⁴³. The interaction of various immune checkpoints with their respective ligands is summarized in Figure 1.

T-CELL ACTIVATION

Two effector cells of adaptive immunity include cytotoxic T cells and helper T cells. Helper T cells contribute to the propagation of the antitumor immune response, and cytotoxic T cells kill tumor cells directly. T-cell activation is necessary for the functioning of both cell types⁷. T-cell activation is essentially a strongly controlled process⁴⁷. For efficient T-cell activation, activation signals must be presented. To explain T-cell activation, a three-signal hypothesis has



Figure 1: Interaction of various immune checkpoints with their respective ligands. The different colors represent different receptors with their specific ligands. APC: Antigen Presenting Cell, ICI: Immune Checkpoint Inhibitor, PD-L: Programmed Death-Ligand, B7-1/2: Immune-regulatory Ligand, MHC: Major Histocompatibility Complex, HVEM: Herpes Virus Entry Mediator, PD-1: Programmed Death-1, CTLA-4: Cytotoxic T-lymphocyte-associated Antigen 4, LAG-3: Lymphocyte-activation Gene 3, TIM-3: T-cell Immunoglobulin and Mucin Domain 3, BTLA: B and T-lymphocyte Attenuator.

been formulated. Signal 1 is given by the interaction between the APCs of MHC and TCR^{20,48}. CD4 and CD8 coreceptors are related to TCR. CD4 binds with MHC II, whereas CD8 binds with MHC I⁴⁹. Next, T-cell surface molecules provide a costimulatory signal that serves as signal 2. CD28 is the most effective and best-characterized costimulatory molecule associated with T-cell activation. This 44-kDa glycoprotein molecule binds to CD80 and CD86 on the APCs. Finally, cytokines act as signal 3 for the activation of T cells. However, CD28 signaling increases cytokine production by T cells and increases T-cell survival by upregulating BCL-X_L, a member of the BCL-2 family of proteins^{20,35,48,50}.

INTERDEPENDENCY OF ICI AND T-CELL ACTIVATION

PD-1 interaction with its ligands alleviates T-cell functions by restraining the effector activity of T cells against cancers^{9,51,52}. Ligand binding with PD-1 causes tyrosine phosphorylation of the PD-1 cytoplasmic domain and recruits both SHP-2 and SHP-1. These events reduce the phosphorylation of TCR signaling molecules and decrease cytokine production and T-cell activation. PD-1 signals can decrease

the expression of anti-apoptotic genes but increase the expression of proapoptotic genes. Additionally, PD-1 signals can also decrease the killing capacity of T cells by reducing their production of cytotoxic molecules^{8,32,40,52,53}. Ligand binding to PD-1 also inhibits the PI3K/AKT pathway and downregulates Bcl-xl gene expression to enhance T-cell apoptosis¹¹. The PD-L2 ligand has a 3-fold increased affinity for PD-1 compared with PD-L1. However, its expression is restricted mainly to myeloid cells. Alternatively, PD-L1 is primarily expressed in nonhematopoietic as well as hematopoietic cells 28,54,55. PD-1 engagement by PD-L2 prevents TCR-mediated proliferation along with cytokine production. Moreover, PD-L2 and PD-1 interaction inhibits B7-CD28 signals at lower antigen concentrations. However, at higher concentrations, these interactions reduce the production of cytokines but do not inhibit T-cell proliferation. These interactions are also found to arrest the cell cycle at the G_0/G_1 phase but do not increase cell death²⁵.

CTLA-4 is a negative controller of the T-cell response, and it prevents the T-cell response in 2 ways^{8,33–35,56}. First, it intrinsically inhibits T-cell activation by outcompeting CD28 to binding with the ligand B7 or reducing TCR and CD28 signaling, recruiting phosphatases to the CTLA-4 cytoplasmic domain. Second, CTLA-4 of one T-cell prevents the activation of another T-cell, lessening the expression of CD80 and CD86 on APCs^{8,53,54}. Again, CTLA-4 can also inhibit TCR signal transduction by binding to the TCR zeta chain and preventing tyrosine phosphorylation⁵⁶.

BLOCKING OF IMMUNE CHECKPOINTS

Immune surveillance employs both the innate and adaptive immune systems and clears early malignant cells in cancer. However, tumors employ various mechanisms to escape this action of immune surveillance. For example, checkpoint blockade by ICIs has dramatically changed cancer treatment by activating the immune system of patients²⁶. Checkpoint blockade in cancer immunotherapy uses antibodies to block the pathways that prevent responses of T cells to tumors⁸.

PD-1 and PD-L1 interaction blockade has exhibited a tremendous antitumor response^{57,58}. PD-1 pathway blockade has exhibited 30%-50% response rates in clinical trials for various cancers⁵⁹. Even more, pembrolizumab, a member of the anti-PD-1 family, has gained the approval of the US Food and Drug Administration (FDA) in melanoma cancer treatment^{8,9}. Furthermore, in 2013, this humanized IgG4- κ monoclonal antibody (mAb), previously known as lambrolizumab and MK-3475, received the "Breakthrough Therapy" designation for advanced melanoma from the FDA^{21,39,60,61}. Pembrolizumab has also been successful in non-small cell lung cancer (NSCLC) treatment⁵⁵. It blocks PD-1 and thus releases it from interactions with ligands and provokes an antitumor response^{40,62}.

Nivolumab (MDX-1106 or BMS-936558), an anti-PD-1 mAb, has also received approval from the FDA for cancers, such as squamous cell lung cancer, renal cell carcinoma, and melanoma treatment^{8,9,40,53,63}. In 2014, this human IgG4- κ mAb achieved the "Breakthrough Therapy" designation from the FDA for non-Hodgkin's lymphoma treatment^{21,61}. Nivolumab blocks the binding of PD-1 with its ligands and thereby augments the host antitumor response by attenuating inhibitory signals^{14,64}. As a result, nivolumab provides significant clinical activity and an acceptable safety profile in squamous NSCLC¹⁴. Additionally, both pembrolizumab and nivolumab prolong overall survival compared to the chemotherapy drug, docetaxel, and have received FDA approval as second-line treatments in advanced NSCLC⁶⁵.

Pidilizumab, an IgG1- κ mAb blocks PD-1, and BMS-936559, an IgG4 mAb, prevents PD-L1 binding with CD80 and PD-1^{21,39}. Again, MPDL-3280A is another IgG1- κ mAb to PD-L1. In its Fc region, it possesses a single amino acid substitution^{16,39,66}. This monoclonal antibody has received approval from the FDA for NSCLC and bladder cancer treatment⁹.

The first clinically targeted immune checkpoint receptor is CTLA-4²⁴. CTLA-4 blockade involves the induction of long-term regression of tumors⁵⁹. Anti-CTLA-4 antibodies increase the ratio of CD4⁺ and CD8⁺effector T cells (Teffs) to forkhead box protein 3 (FoxP3)⁺Tregs in tumor infiltrates, which is associated with the enhanced eradication of tumors^{8,55,67}. An increased ratio of Teffs to Tregs is a hallmark of successful tumor rejection with CTLA-4 blockade⁶⁸. Recent studies suggest that the therapeutic outcome of anti-CTLA-4 antibodies is possibly not just because of the CTLA-4 interaction blockade with ligands but also because of the lessening of intratumoral Tregs through Fc receptor-mediated cytotoxicity^{8,66}.

The first FDA-approved ICI was ipilimumab (MDX-010). This human IgG1- κ mAb blocks CTLA-4. The half-life of ipilimumab is 15.4 days⁶⁶. Ipilimumab received FDA approval in 2011 for metastatic melanoma treatment^{9,39,61}. It has been assessed in clinical trials with renal cell carcinoma, prostate cancer, and melanoma malignancies²¹. Additionally, this was the first ICI to be evaluated in NSCLC⁶⁶. In addition, it has improved overall survival in advanced cutaneous melanoma¹⁵. It blocks the CTLA-4 receptor from interacting with the ligands B7-1 and B7-2^{7,55,64}. It also inhibits the signaling activity of CTLA-4⁶¹. Tremelimumab, a humanized IgG2 mAb blocking CTLA-4, has also entered clinical trials^{21,39,40,53,66}.

Various anti-LAG-3 antibodies are now being evaluated in clinical trials, with relatlimab being the first anti-LAG-3 antibody developed commercially³⁸. Blocking LAG-3 inhibits Tregs and restores cytotoxic T-cell function²⁰. Zhou *et al.* identified LBL-007 and reported that this novel humanized anti-LAG-3 antibody demonstrated a higher affinity for LAG-3 antigen and blocked downstream signaling and functions of LAG-3. They also reported that LBL-007 suppressed colorectal cancer cell growth in mice when used alone or with a PD-1 antibody³⁸.

Unfortunately, the blocking mechanism of TIM-3 mAbs is not yet fully understood⁸. However, it has been found that targeting the Gal9/TIM-3 axis together with induction chemotherapy could effectively

increase the possibility of total remission of acute myeloid leukemia. Again, combination therapy of PD-1 and TIM-3 mAb can regulate tumor growth in a synergistic way⁶⁹. In melanoma and mammary carcinoma models, anti-BTLA antibodies have demonstrated significant antitumor activity⁴⁴. Further, it has been found that the duel blockade of BTLA and PD-1 enhances antitumor immunity²⁴ and releases immune exhaustion in tumor-specific T cells in melanoma patients. Hence, BTLA inhibition with other blocking antibodies is a possible therapeutic approach in cancer immunotherapy⁴⁵. A list of the bestknown ICIs for immunotherapy is presented in **Table 1**.

COMBINATION THERAPY

Combination therapy helps enhance various steps in the cancer-immunity cycle to establish an active microenvironment where ICIs can exert successful antitumor killing⁸. As ICIs are becoming a mainstream treatment option in cancer, different combinations of various ICIs or ICIs with other proven treatment options are being tested to convert nonresponders to responders. PD-1 blockade-associated toxicity is less than CTLA-4 and IL-2 blockade. However, PD-1 and CTLA-4 combination checkpoint blockades improve the response rate compared with single checkpoint blockades^{8,9}. Furthermore, a CTLA-4 and PD-1 combination checkpoint blockade allows tumorspecific T cells to accomplish effector functions and increases the infiltration of Teff. So, it increases the advantageous ratio of Teff-to-Tregs and causes the tumor-suppressive microenvironment to shift to the inflammatory microenvironment^{51,61}. Again, LAG-3 and PD-1 combination blockade also exhibited increased antitumor activity compared with their single checkpoint blockade^{8,38}. Nivolumab and ipilimumab combination therapy has proven to be the most active immunotherapy regarding response rate, along with median progression-free survival in melanoma treatment^{40,55,70}. The combination of nivolumab and relatlimab is being assessed for immunotherapy in solid cancers³⁸.

Cancer therapies, such as radiotherapy, oncogenetargeted therapy, or chemotherapy, can change the immunosuppressive tumor microenvironment and synergize with ICIs. Thus, various combinations of ICIs with other cancer therapies are being tested⁵³. For example, clinical trials have assessed ipilimumab and radiotherapy⁵⁴. Golden EB *et al.* reported the first abscopal response (radiotherapy-induced regression of tumors in lesions distant from a targeted site) with the combination of radiotherapy and ipilimumab in lung cancer patients. They reported that irradiating single liver metastasis in a lung cancer patient with ipilimumab led to a durable and complete response^{54,68}. Therefore, radiation therapy can emerge as an optimal partner for ICIs because of its ability to induce a response in nonresponsive patients. For many years, chemotherapeutic agents have been used to control cancers that do not respond to either radiation or surgery⁶. In addition, chemotherapeutic agents can initiate T-cell activation and sensitize tumor cells for T-cell-mediated killing by inducing tumor-specific antigen release. These observations inspired combining chemotherapeutic agents with immunotherapy to enhance patient response rate³⁹. As ipilimumab is an FDA-approved ICI, the possible role of ipilimumab in combination with chemotherapy, other targeted therapies, and immunotherapy is being evaluated 40. Thus, ipilimumab has been evaluated as a combination therapy with chemotherapeutic agents in small cell lung cancer (SCLC) and NSCLC in clinical trials^{66,68}. When used alone, ipilimumab has virtually no effect on lung cancer. However, combined with chemotherapeutic agents, it seems to provide a modest benefit in NSCLC and SCLC⁷¹.

Small RNAs without coding potential are known as microRNAs or miRNAs. A network of microRNAs regulates the expression of immune checkpoints, directly or indirectly. MicroRNAs can target multiple immune checkpoints and mimic the therapeutic benefit of combined checkpoint blockade. Dragomir *et al.* hypothesized that microRNA combined with a checkpoint blockade could increase the efficacy of established monotherapies⁷².

TOXICITY

The hallmark toxicity associated with ICIs includes immune-related adverse events (irAEs) from aberrant activation of autoreactive T cells against host tissue. Substantial morbidity or even mortality may be infrequently caused by irAEs^{22,73}. However, these toxicities differ significantly from conventional chemotherapeutic toxicities⁷⁴. Multiple organs, such as the peripheral and central nervous system, kidney, liver, pancreas, eyes, and skin, can be affected by irAEs^{32,75}. Moreover, immune toxicities, such as dermatitis, pneumonitis, diarrhea, colitis, thyroiditis, and hepatitis, are very common with ICIs. Pneumonitis is very rare but a potential life-threatening irAE of CTLA-4 and PD-1 inhibitors^{26,76,77}. In monotherapy, thyroid disorders are almost the same for CTLA-4 and PD-1 inhibitors but remarkably increase in combination

Immune Checkpoint Inhibitor (ICI)	Target Immune Checkpoint	Antibody Isotype	Appoval Status	References
Pembrolizumab	PD-1	Ig G4	FDA-approved	8,9,21,39,60,65
Nivolumab	PD-1	Ig G4	FDA-approved	8,9,21,40,53,61,63
MPDL-3280A	PD-L1	Ig G1	FDA-approved	9,16,39,66
Ipilimumab	CTLA-4	Ig G1	FDA-approved	9,39,61
Tremelimumab	CTLA-4	Ig G2	Under Assessment	21,39,40,53,66

Table 1: List of the best known Immune Checkpoint Inhibitors (ICIs) for Immunotherapy

therapy. Symptoms of thyroid disorders include fatigue, alopecia, constipation, and palpitations⁷⁸.

In general, PD-1 inhibitors demonstrate fewer irAEs than CTLA-4 inhibitors^{76,79}. However, thyroid dysfunction is higher in cases of anti-PD-1 antibodies (e.g., pembrolizumab and nivolumab)^{80,81}. Commonly reported adverse events of PD-1 inhibitors include rash, pruritus, fatigue, constipation, diarrhea, and arthralgia. Dry mouth and oral mucositis are more frequent with PD-1 inhibitors⁷⁴. Pneumonitis is a very concerning toxicity associated with nivolumab⁶⁶. On the other hand, hypophysitis has been reported as a characteristic side effect associated with CTLA-4 inhibitors. If not promptly identified, this situation can be life-threatening owing to secondary adrenal insufficiency^{15,78}. Common reported irAEs of ipilimumab include hepatotoxicity, diarrhea, endocrinopathies, and dermatologic toxicity⁷⁴. Again, hypophysitis is frequently reported with ipilimumab⁸⁰. Thyroid dysfunction is very common for pembrolizumab, but hypophysitis is found to be infrequent. Again, thyroid disorders are frequently reported irAEs for tremelimumab⁸². It has been found that irAEs with CTLA-4 blockade increase as the dose increases. However, irAEs with PD-1 blockade are not related to the dose^{7,79,82}.

Toxicity associated with combination therapy is a matter of great concern⁷⁶. Colitis is severe and more frequent with combination therapy⁸³. Combination therapy of ipilimumab and nivolumab has demonstrated higher irAEs than their use in monotherapy. For combination therapy with ipilimumab with nivolumab, rash is the most commonly reported toxic effect, and this combination shows more frequent thyroid dysfunction and sometimes a higher grade^{76,80}.

BIOMARKER

PD-1 or PD-L1 monotherapy is considered well tolerated, but irAEs increase in combination therapy. Therefore, the development of biomarkers can contribute to maximizing therapeutic benefit, minimizing irAEs, and guiding combination therapy⁸⁴. Predictive biomarkers help to determine the result of therapy before starting it. These can indicate whether a patient would benefit from monotherapy or if there is a need for combination therapy⁸⁵. PD-1 is expressed on the activated T-cell surface. Therefore, it can be used as an activation marker⁸⁶. Again, PD-L1 expression on specimens of pretreatment tumors can be a predictive biomarker with inhibitors of PD-1^{39,84,87}. Detection of PD-L1 with immunohistochemistry is thus far a commonly detected clinical biomarker to predict anti-PD-1/PD-L1 therapy response⁸⁶.

Again, the status of baseline TILs and the mutational or neoantigen burden have been evaluated as predictive biomarkers in immunotherapy. In addition, peripheral blood marker testing is a noninvasive source of possible biomarkers in ICI therapy. However, no predictive peripheral blood marker has yet been validated ⁸⁴.

CONCLUSION

ICIs offer a blossoming field in cancer treatment. It has already shown valuable clinical benefits in several tumor types. Again, combination blockade by checkpoint inhibitors can exert better patient response rates. Therefore, a combination blockade of different checkpoint inhibitors needs to be tested. Although ICIs show less toxicity than previous immunotherapies, further study is required for a complete understanding of the side effects of these treatments. Further challenges in ICI therapy include exploring new therapeutic targets, optimizing dosing regimens, and identifying and validating biomarkers for predicting toxicity and clinical responses. Identifying reliable biomarkers will help make better treatment decisions regarding efficacy and toxicity. We hope the current manuscript will help us understand the basic principle of ICI therapy and its prospects in cancer treatment.

ABBREVIATIONS

aa: Amino Acid, APC: Antigen-presenting Cell, BTLA: B- and T-lymphocyte Attenuator, CTLA-4: Cytotoxic T-lymphocyte-associated Antigen 4, DC: Dendritic Cell, FDA: US Food and Drug Administration, FoxP3: Forkhead Box Pprotein 3, HVEM: Herpes Virus Entry Mediator, ICI: Immune Checkpoint Inhibitor, Ig: Immunoglobulin, IL: Interleukin, irAEs: Immune-related Adverse Events, ITIM: Immunoreceptor Tyrosine-based Inhibitory Motif, ITSM: Immunoreceptor Tyrosinebased Switch Motif, LAG-3: Lymphocyte-activation Gene 3, mAb: Monoclonal Antibody, MHC: Major Histocompatibility Complex, NK: Natural Killer, NSCLC: Non-small Cell Lung Cancer, PD-1: Programmed Death 1, SCLC: Small Cell Lung Cancer, SH2: Src Homology 2, SHP-1: SH2-containing Tyrosine Phosphatase 1, SHP-2: SH2-containing Tyrosine Phosphatase 2, TCR: T-cell Receptor, Teffs: Effector T Cells, TIL: Tumor-infiltrating Lymphocyte, TIM-3: T-cell Immunoglobulin and Mucin Domain 3, Tregs: T-regulatory Cells

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The authors declare that they have no competing interests.

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