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Targeting the Programmed Death-1 Pathway in Lymphoid Neoplasms

Dr. Chi Young Ok, MD1 and **Dr. Ken H. Young, MD, PhD**¹

¹The Department of Hematopathology, The University of Texas MD Anderson Cancer Center, Houston, Texas

Abstract

Programmed death-1 (PD-1) is a co-inhibitory molecule and is seen in CD4+ and CD8+ T cells. Upon binding to its ligands, programmed death ligand-1 (PD-L1) and -2 (PD-L2), PD-1 negatively regulates interleukin 2 (IL-2) production and T cell proliferation. Activated effector T-cells, which kill cancer cells, can be affected by PD-1 signaling in some lymphoid neoplasm that express PD-L1 or PD-L2. PD-L1 expression in tumor cells can be induced by extrinsic signal (i.e. interferon gamma) or intrinsic signals, such as genetic aberrations involving 9p24.1, latent Epstein-Barr virus infection, PD-L1 3′-untranslated region disruptions, and activated Janus kinase/signal transducer and activator of transcription (JAK/STAT) pathway. Anti-PD-1 therapy improves the overall response rate to treatment in patients with lymphoid neoplasms, particularly relapsed/refractory classical Hodgkin lymphoma. Inspired by their success in treating patients with classical Hodgkin lymphoma, medical practitioners have expanded PD-1 therapy, given as a single therapy or in combination with other drugs, to patients with other types of lymphoma. In this review, current clinical trials with anti-PD-1 or anti-PD-L1 drugs are summarized. The results of numerous clinical trials will broaden our understanding of PD-1 pathway and shall expand the list of patients who will get benefit from these agents including those who suffer from lymphoid neoplasms.

Graphical Abstract

Correspondence: Dr. Chi Young Ok, MD, The University of Texas MD Anderson Cancer Center, Department of Hematopathology, 1515 Holcombe Boulevard, Houston, TX 77030, USA. Office: (713)792-7030, COk@mdanderson.org. Dr. Ken H. Young, MD, PhD, The University of Texas MD Anderson Cancer Center, Department of Hematopathology, 1515 Holcombe Boulevard, Houston, Texas 77230-1439, USA. Phone: 713-745-2598, khyoung@mdanderson.org.

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Keywords

lymphoid neoplasms; PD-1; PD-L1; PD-L2; immune checkpoint

Introduction

T cell activation is explained by the two-signal model proposed by Bretscher and Cohn (Figure 1).[1] The first signal, which triggers T cell receptor (TCR) signaling, occurs when the major histocompatibility complex (MHC) molecule of an antigen-presenting cell (APC) presents a processed antigen to the TCR of a T cell. The second signal, which is an antigenindependent, co-stimulatory or co-inhibitory signal delivered by the APCs, modulates TCR signaling and determines the T cell's fate. The prototypical molecule of the second signal is CD28, which constitutively expressed in resting naïve T-cells.[2] CD28's ligands (such as B7-1 or B7-2) induce cell-cycle progression, interleukin-2 (IL-2) production, and clonal expansion. If T cells do not receive co-stimulatory second signals from molecules like CD28, they become anergic. In contrast, cytotoxic T-lymphocyte antigen-4 (CTLA-4), which shares the same ligands with CD28, is a co-inhibitory receptor on T-cells that induces T-cell tolerance.[3] If T cells do not receive co-inhibitory second signals, fatal lymphoproliferation and multiorgan autoimmunity can occur in mice.[4] Additional secondsignal receptors and ligands, which are collectively called B7-CD28 family, have also been discovered and include programmed death ligand-1 (PD-L1) and PD-L2. Unlike B7-1 and B7-2, PD-L1 and PD-L2 do not bind to CD28 or CTLA-4. Instead, they bind to programmed death-1 (PD-1) on T-cells.

The PD-1 pathway has recently emerged as an attractive target in cancer immunotherapy. In the context of the cancer-immunity cycle, the aim of immunotherapy is to restore immune function at various steps of cancer-immunity cycle. Several clinical studies have shown that blocking the PD-1 pathway leads to significant responses in patients with various solid tumors.[5] Numerous clinical trials with PD-1 pathway blocking agents, used either alone or in combination with other therapies, are currently in progress. In this review, we discuss the functions of the PD-1 pathway in the cancer-immunity cycle, the armamentarium of PD-1

pathway blocking agents, current and prior clinical trials in patients with lymphoid malignancies and future directions in search.

The structure of PD-1, PD-L1, and PD-L2

PD-1 is a protein encoded by the 5-exon PDCD1 gene on chromosome 2q37.3. It consists of 288 amino acids, and its calculated molecular weight is 31.6 kDa. However, Agata et al's [6] immunoprecipitation of the protein revealed broad bands with molecular weights of 50–55 kDa, suggesting that the protein is heavily glycosylated. PD-1 contains a single immunoglobulin V-like domain, a transmembrane domain, and an intracellular domain with an immunoreceptor tyrosine-based inhibitory motif (ITIM) and an immunoreceptor tyrosinebased switch motif (ITSM).[7,8]

Two PD-1ligands, PD-L1 and PD-L2, have structures similar to that of PD-1 in that they contain an immunoglobulin V-like domain, an immunoglobulin C-like domain, a transmembrane domain, and an intracellular domain.[9] These ligands interact with PD-1 via their immunoglobulin V-like domains. PD-L1 is encoded by the 8-exon CD274 gene on chromosome 9p24.1, is composed of 290 amino acids, and has a molecular weight of 33.3 kDa. Of note, it also competitively binds to B7-1, thus inhibiting the CD28-mediated costimulation of T-cells.[10]

PD-L2 is encoded by the 7-exon PDCD1LG2 gene, and is located 42 kilobases apart from the CD274 gene on chromosome 9p24.1. It consists of 273 amino acids, and its molecular weight is 31.0 kDa.[11]

The PD-1 signaling pathway

PD-1 is present on T-cells as a monomer and is a negative regulator of IL-2 production and T-cell proliferation.[12,13] PD-1 inhibition of antigen receptor signaling is only seen when PD-1 ligation occurs close to the site of antigen receptor engagement.[13] Indeed, it has been observed that randomly located PD-1 migrates to the immunological synapse during the interaction between T cell and APC.[14] Once PD-1 has bound to ligands, its ITIMs and ITSMs are phosphorylated by the Src-family tyrosine kinases (Figure 2). The phosphorylated tyrosine residue subsequently recruits Src homology 2 domain-containing phosphatases (SHPs), which dephosphorylate signaling intermediates and down-regulate TCR signaling. Of note, ITIM recruits only SHP-2, but ITSM recruits both SHP-1 and SHP-2.[15, 16] SHP-2 appears to be more important than SHP-1 in PD-1 signaling because T cell stimulation with PD-L2 increases the amount of SHP-2 but not of SHP-1.[17] In addition, ITSM is more important than ITIM in PD-1 signaling because PD-1's inhibitory function is lost when ITSM is mutated but not when ITIM is mutated.[15, 18, 19]

PD-1 inhibits phosphatidylinositol 3-kinase (PI3K)/Akt pathway by thwarting CD28 mediated activation of PI3K via ITSM. In contrast, CTLA-4 bypasses PI3K and instead halts Akt induction via the intracellular serine/threonine phosphatase PP2A.[20] PD-1 can also block the RAS/MEK/Erk pathway. Of interest, PD-1 inhibits the PI3K/Akt pathway within minutes, whereas it takes a few hours for it to block the RAS/MEK/Erk pathway.[17] Ultimately, PD-1's inhibition of both of these pathways halts cell cycle progression.[21] In

addition, PD-1's inhibition of the PI3K/Akt pathway prevents T cell's expression of the antiapoptotic protein Bcl-xL, which depends upon PI3K.[20] PD-1 also hinders phosphorylation of ZAP70, an essential molecule for T-cell activation; inhibits activation of PKC-θ, which is critical for IL-2 production, cell cycle progression and T-cell activation; and prevents effector T-cell development by inhibiting glycolysis and promoting fatty acid oxidation.[16, 22, 23] Of note, PD-1 mediated inhibitory signals are inversely associated with the strength of the TCR signal. Furthermore, PD-1 inhibition can be overcome by T cell stimulation with CD28 or exogenous IL-2.[24]

The PD-1 pathway plays an important role in enabling tumor cells to evade the immune response. The rate of tumor lysis by cytotoxic T cells (CTLs) in vitro was lower in P815 murine mastocytoma tumor cells with transgenic expression of PD-L1 than in parental tumor cells without PD-L1 expression. Furthermore, inoculating syngenic BALB/c mice with PD-L1 expressing myeloma cells led to more rapid tumor growth, an outcome which was then suppressed by anti-PD-L1 antibodies.[25] Similarly, another study with a mouse model showed that PD-L1 expression in tumor cells confers resistance to activated CTLs but that this effect could be overcome with anti-PD-L1 or anti-PD1 antibodies.[26] In addition, Dong et al's [27] study showed that PD-L1 expression in a melanoma cell line (624mel) enhanced apoptosis of PD-1-expressing CTLs.

Expression of PD-1, PD-L1, and PD-L2 in normal tissue

PD-1 is expressed in activated CD4+ and CD8+ T cells, naïve and activated B cells, myeloid dendritic cells, and with low intensity in monocytes. It is not expressed in resting T cells, but its expression can be induced within 24 hours of T cell activation.[28] In normal human tissue, PD-L1 expression is seen on follicular T-cells, macrophages and a subset of dendritic cells in lymphoid tissue, placental syncytiotrophoblasts, and dendritic cells/monocytes in the lung and liver, although low levels of PD-L1 mRNA are seen in almost all normal tissue. [28–30] Expression of PD-L2 is more restricted compared with PD-L1. Low levels of PD-L2 expression are seen in activated CD4+ or CD8+ T cell subsets, myeloid dendritic cells, monocytes, endothelial cells, and placental syncytiotrophoblasts.[31] Expression of PD-L1 and PD-L2 is induced by inflammatory signals such as interferon gamma (IFN- γ), granulocyte macrophage colony-stimulating factor (GM-CSF), and IL-4.[27, 32–34]

Expression of PD-L1 in lymphoid malignancies

In contrast to the infrequent PD-L1 expression in normal tissue, immunohistochemistry shows that expression of PD-L1 is often found in a variety of cancers.[5] However, the expression pattern of PD-L1 in tumor cells is not uniform. In some tumors, heterogeneous expression of PD-L1 is seen at the interface of the tumor and tumor-infiltrating lymphocytes (TILs).[35] In such cases, PD-L1 expression is likely to be induced by IFN-γ secreted from the TILs.[36] Homogeneous expression of PD-L1 is observed in other tumors, particularly in the Reed-Sternberg (RS) cells of classical Hodgkin lymphoma (CHL).[37] Homogeneous PD-L1 expression usually correlated with intrinsic signals, which are reviewed below. The underlying mechanisms of PD-L1 expression are different in different types of cancer, and antibodies targeting different domains of PD-L1 have been used in various studies (Table 1).

[35, 37, 38] Furthermore, companion diagnostic assays were developed specifically for trials of particular anti-PD1 or anti-PD-L1 agents, including 22C3 for pembrolizumab, 28-8 for nivolumab, and SP142 for atezolizumab.[35, 39, 40] Therefore, PD-1 pathway investigators should pay attention to antibody types when designing a study or interpreting clinical trial results.

As determined by immunohistochemistry, PD-L1 expression is present in CHL; in 87% of CHL cases, irrespective of subtype, over half of RS cells are PD-L1-positive.[41] A recent study demonstrated that almost all (97%) patients with CHL harbored either polysomy, copy-number gain or amplification of 9p24.1 detected by fluorescence in situ hybridization (FISH).[42] In approximately 40% of CHL cases of the nodular sclerosis subtype, the RS cells have copy number alterations (amplifications or gains) and/or translocations involving 9p24.1/PD-L1/PD-L2.[43] Furthermore, Green and colleagues demonstrated that amplification of 9p24.1 not only increases the genetic dosage of PD-L1/PD-L2 but also induces JAK2 amplification and, consequently, enhancement of JAK/STAT signaling.[43] Since PD-L1 has a promoter which is responsive to Janus kinase/signal transducer and activator of transcription (JAK/STAT) signaling pathway, extra signaling for PD-L1 expression is present in CHL. Advanced stage (stages III/IV) CHL patients with 9p24.1 amplification have significantly shorter progression-free survival.[42]

Copy number alterations and/or translocations involving 9p24.1/PD-L1/PD-L2 are also frequently seen in patients with primary mediastinal large B-cell lymphoma (PMBL) (~70%), EBV-negative primary central nervous system lymphoma (PCNSL) (~60%), and primary testicular lymphoma (PTL) $({\sim}60\%)$.[43, 44, 45] Immunohistochemistry shows that approximately 70%, 50% and 50% of PMBL, PCNSL and PTL tumors, respectively, express PD-L1, supporting the idea that PD-L1 expression is induced by cytogenetic abnormalities. [41, 44]

In DLBCL, not otherwise specified (DLBCL, NOS), PD-L1 expression is seen in 11–26% of cases by immunohistochemistry.[41, 46, 47] Interestingly, a similar percentage $(\sim 20\%)$ of cytogenetic abnormalities in 9p24.1 (gains, amplifications or translocations) is also observed in these cases.[43, 46, 47] Gains and amplifications of 9p24.1 and translocations of PD-L1 correlate with increased PD-L1 mRNA expression and PD-L1 protein expression. Furthermore, cytogenetic abnormalities in 9p24.1 and PD-L1 protein expression are associated with activated B-cell-like or non-germinal center B cell-like phenotype of DLBCL, as determined by gene expression profiling and Hans classification, respectively. [46, 47] A Japanese study determined that PD-L1 expression in DLBCL is an independent indicator of poor prognosis.[47] Available data suggest that, in DLBCL patients, the PD-1 pathway could be an immune escape mechanism and that patients with PD-1 expression could be good candidates for anti-PD-1 or anti-PD-L1 therapies.

EBV provides an intrinsic signal to augment PD-L1 expression. EBV latent membrane protein 1 (LMP1) activates the JAK/STAT pathway and the transcription factor AP-1; this enables JAK3 to activate a PD-L1 promoter and AP-1 to stimulate a PD-L1 enhancer.[48– 50] Chen et al. [41] have shown that PD-L1 expression in RS cells is more commonly present in EBV-positive CHL than in EBV-negative CHL. Similarly, PD-L1 expression is

seen in all cases of EBV-positive diffuse large B-cell lymphoma (EBV+ DLBCL) and EBVpositive immunodeficiency-related DLBCL.[41] Unlike EBV-negative PCNSL, EBVpositive PCNSL frequently express PD-L1 without 9p24.1/*PD-L1*/*PD-L2* copy number alterations, suggesting that PD-L1 expression is induced by viral proteins.[44] Other EBVassociated lymphoproliferative disorders including EBV+ post-transplant lymphoproliferative disorder (PTLD), plasmablastic lymphoma, primary effusion lymphoma, and extranodal NK-T-cell lymphoma express PD-L1 in approximately 60%, 50%, 50% and 70%, respectively.[41, 50]

Chen and colleagues showed that 91% of T-cell rich, histiocyte-rich large B-cell lymphomas (TCHRBCLs) express PD-L1, compared to only 13% of nodular lymphocyte predominant Hodgkin lymphomas (NLPHLs).[41] Since TCHRBCL and NLPHL have considerable morphologic overlap but opposite clinical behaviors, this finding is very intriguing.[51] One can speculate that acquiring PD-L1 expression is an immune escape mechanism of indolent NLPHL and that leads to more aggressive TCHRBCL. Of note, histiocytes adjacent to lymphoma cells also show strong PD-L1 expression in TCHRBCL, suggesting that both tumor cells and background inflammatory cells provide immune escape signals.[41] Since 9p24.1 aberrations and EBV infection are not generally seen in TCHRBCL, PD-L1 is thought to be induced by IFN- γ secreted from tumor-infiltrating T-cells.[52]

Another intrinsic mechanism of increased PD-L1 expression was revealed by Kataoka and colleagues, who showed that increased PD-L1 expression is associated with PD-L1 3′ untranslated region (UTR) disruption in about 30% and 10% of adult T-cell leukemia/ lymphoma patients and DLBCL patients, respectively.[53] Of interest, PD-L1 expression induced by 3′-UTR disruption with truncated protein was only seen when an antibody was directed against the extracellular domain and not when an antibody was directed against the cytoplasmic domain. Whether 3'-UTR disruption in *PD-L1* is present in other lymphoid neoplasms is still unknown.

PD-L1 expression also occurs through constitutive activation of the JAK/STAT pathway. In ALK+ anaplastic large cell lymphoma with NPM-ALK rearrangement, the fusion transcript has been shown to induce expression of PD-L1 via STAT3 activation.[54] Furthermore, PD-L1 expression in DLBCL correlates the ABC phenotype, which is known for its enhanced JAK/STAT pathway.[55]

PD-L1 expression in myeloma cells can be detected by flow cytometry.[56] Significant increase in copy number of PD-L1 and PD-L1 mRNA expression is seen in malignant myeloma cells, which correlates with PD-L1 expression in myeloma cells.[56] Compared with normal plasma cells, PD-L1 expression is up-regulated in non-hyperdiploid and hyperdiploid myeloma cells, with higher expression in the latter. However, PD-L1 expression is not associated with del (1q), del (13), del(17p), $t(11;14)$, $t(4;14)$ or $t(14;16)$.

PD-L1 expression is not generally seen in follicular lymphoma, mantle cell lymphoma, marginal zone lymphoma, Burkitt lymphoma or chronic lymphocytic leukemia/small lymphocytic lymphoma.[57, 58]

Expression of PD-L2 in lymphoid malignancies

Lymphoid neoplasms with abnormalities in 9p24.1/*PD-L1*/*PD-L2* generally express PD-L2. RS cells in CHL also express PD-L2, irrespective of EBV status.[59] PD-L2 expression by immunohistochemistry occurs in 72% of PMBL, and copy number gain of *PDCD1LG2* gene has also been observed in most cases.[60] PD-L2 expression is also present in PTLs and PCNSLs, which are associated with 9p24.1/PD-L1/PD-L2 copy gains or translocations.[44] However, unlike PD-L1 expression, PD-L2's expression of RNA and protein were not associated with cytogenetic abnormalities in 9p24.1 in DLBCL, NOS.[46] Also, PD-L2 expression is not associated with EBV infection or $3'$ -UTR disruption in the *PD-L1* gene. [53, 57]

Expression of PD-1 in the lymphoid malignancy microenvironment

Although PD-1 expression is seen in a few lymphoid malignancies, particularly T cell lymphomas with the follicular helper T cell phenotype, most lymphoid neoplasms do not express PD-1.[30, 61] Thus, PD-1 expression is best examined in the microenvironment. PD-1 expression in TILs has been reported in follicular lymphoma, and NLPHL is well known for PD-1-expressing T-cell rosettes surrounding neoplastic L&H cells (LP cells or popcorn cells).[62, 63] Since both neoplasms arise from germinal center B cells, it is not surprising that their microenvironments are like those of their normal counterparts. Likewise, PD-1-expressing TILs are also correlated with DLBCL, germinal center B-celllike (GCB) phenotype.[47] Interestingly, the high number of PD-1-expressing TILs in the DLBCL microenvironment is inversely associated with PD-L1 expression in lymphoma cells. Furthermore, contrary to observations in solid tumors, the high number of PD-1 expressing TILs in DLBCL and in follicular lymphoma (FL) is associated with a favorable prognosis; this suggests that the presence of PD-1-expressing TILs in lymphomas could simply indicate cell-of-origin, unlike tumor-mediated T-cell exhaustion in solid tumors.[47, 62, 64, 65]

Anti-PD-1 antibodies

Pembrolizumab (Keytruda®, MK-3475, SCH 900475, previously lambrolizumab)

Pembrolizumab is a fully humanized IgG4 kappa isotype anti-PD-1 monoclonal antibody. It is well tolerated and is associated with durable anti-tumor activity in multiple solid tumors. [66] In a phase 1b study (KEYNOTE-013 study, NCT01953692), pembrolizumab was administered to 31 patients with CHL every 2 weeks (10 mg/kg IV) until disease progression.[67] The median age was 32 years (range, 20–67 years). Seventeen (55%) patients had ≥ 5 lines of prior therapy, 31 (100%) failed with brentuximab vedotin (BV) treatment, 22 (71%) underwent prior autologous stem cell transplant (ASCT), and 8 (26%) were ineligible for transplantation. Examination of formalin-fixed, paraffin-embedded tissue revealed PD-L1 (clone 22C3) and PD-L2 (clone 3G2) expression in 94% and 90% of tested patients, respectively, before pembrolizumab treatment. With the median follow-up of 17.6 months (range, 10.6–22.5 months), the overall response rate (ORR) was 65%, with 5 (16%) and 15 (48%) patients achieving complete remission (CR) and partial remission (PR), respectively. Among the 20 patients, 16 (80%) achieved their best response around 12th

week of therapy and 14 (70%) had 24 weeks of duration. The progression-free survival (PFS) rates were 69% and 46% at 24 weeks and 52 weeks, respectively. By flow cytometry, expansion of circulating total T cells, including CD4+ and CD8+ T cells, and NK cells was observed at cycle 7 compared to baseline. RNA profiling showed significant increases in the IFN-γ-induced, expanded immune-related, and TCR signatures.[67]

A multicohort phase 2 study is underway (KEYNOTE-087, NCT02453594) to confirm the clinical activity of pembrolizumab in CHL patients. Patients, all of whom have relapsed/ refractory CHL, are in 3 cohorts of those who: (1) had an ASCT and subsequent BV therapy, (2) are ineligible for ASCT due to unresponsiveness to salvage chemotherapy and BV failure, or (3) had an ASCT without BV therapy. At the time of writing, interim analysis of cohorts 1 ($n=30$; median age, 36 years) and 2 ($n=30$; median age, 33 years) has been reported. Similar to the phase 1b study, 40 (67%) patients had ≥ 4 lines of prior therapies. In cohort 1, the ORR was 70%, with 6 (20%) and 15 (50%) patients achieving CR and PR, respectively. In cohort 2, the ORR was 80%, with 8 (27%) and 16 (53%) patients achieving CR and PR, respectively.[68]

A phase 1 study of pembrolizumab given in combination with lenalidomide and low-dose dexamethasone to patients with relapsed/refractory plasma cell myeloma determined that the maximum tolerated dose/maximum administered dose (MTD/MAD) was a 200-mg fixed dose of pembrolizumab combined with 25 mg lenalidomide and low-dose dexamethasone. In the dose determination/confirmation phase, 17 patients were evaluated. The ORR was 76% (13 patients), including 4 patients with a very good PR and 9 patients with a PR. The median duration of response was 9.7 months (range: 0–16.7 months). Updated efficacy data, including data for additional 33 patients in the expansion phase, are forthcoming (KEYNOTE-023, NCT02036502).[69, 70]

Table 2 lists numerous ongoing clinical trials of pembrolizumab given as a single therapy or in combination with other therapies.

Nivolumab (Opdivo®, BMS-936558, MDX-1106, ONO-4538)

Nivolumab is a fully human IgG4 anti-PD-1 monoclonal antibody. It has a high affinity for PD-1 and blocks it from binding to its ligands. Like pembrolizumab, it is well tolerated and is associated with durable anti-tumor activity in solid tumors.[71] In a phase 1b study, 23 patients with relapsed/refractory CHL in whom ASCT and BV was unsuccessful, received 3 mg/kg nivolumab at week 1, week 4, and then every 2 weeks until disease progression or CR or for a maximum of 2 years (NCT01592370).[37] The median age was 35 years (range, 20– 54 years). Twenty (87%) patients had received α 3 lines of prior therapies, 18 (78%) had received BV, and 18 (78%) had undergone ASCT. Ten (43%) patients were tested for PD-L1 and PD-L2 with FISH, and all had polysomy 9p and gain or amplification of PD-L1/PD-L2. Expression of PD-L1 (clone 405.9A11) and PD-L2 (366C.9E5) was seen in all 10 tested patients. At the median follow-up of 40 weeks (range, 0–75 weeks), the ORR was 87%, with 4 (17%) and 16 (70%) patients achieving CR and PR, respectively. Twelve (60%) patients achieved CR or PR by 8 weeks. The PFS rate was 86% at 24 weeks. The median overall survival duration had not been reached.[37]

A subsequent phase 2 study (CHECKMATE 205 cohort B, NCT02181738) recruited 80 patients with relapsed/refractory CHL who received nivolumab (3 mg/kg IV) every 2 weeks. Independent radiologic review committee (IRRC)-determined ORR was 66%, including CR and PR rates of 8.8% and 57.5%, respectively. The investigator-determined ORR was 73%, including CR and PR of 27.5% and 45%, respectively. At the median follow-up 8.9 months, the IRRC 6-month overall survival and PFS rates were 99% and 77%, respectively. Notably, 43 patients with no prior BV response showed an IRRC ORR of 72% with nivolumab treatment.[72]

A Japanese phase 2 study gave 17 patients with relapsed/refractory CHL nivolumab (3 mg/kg) on day 1 of a 14-day cycle. All patients had previously been treated with BV. The ORR in efficacy-evaluable patients was 75%, including 4 (25%) and 8 (50%) patients with CR and PR, respectively. Among 8 BV-resistant patients in the study, 3 had a CR, 4 had a PR and 1 was not evaluable at that time point were observed (JapicCTI-142755).[73]

In a phase 1 study, 81 patients with B-cell lymphoma $(n=31)$, including DLBCL $[n=11]$, and FL [n=10]), T cell lymphoma (n=23), and plasma cell myeloma (n=27) were treated with nivolumab alone (NCT01592370). All patients had received prior systemic treatment regimens (median 3; range, $1-12$). Among the 11 patients with DLBCL, the ORR was 36% , 2 of these patients had a CR, and 2 had a PR. At the median follow-up duration of 22.7 weeks, individual response durations were 6 and 77.3+ weeks for the patients with CR and 12.1+ and 22.1 weeks for the patients with PR, respectively. The ORR of patients with FL was 40%, including 1 patient with CR and 3 patients with PR. At the median follow-up duration of 91.4 weeks, individual response durations were 81.6+ weeks for the patient with CR and 27.1+, 28.1+, and 32.1+ weeks for the patients with PR. Objective responses were not seen in the 10 patients with other B-cell lymphomas. Among the 23 patients with T cell lymphoma, the response rate was 17% (n=4), and all responding patients had PR. Individual response durations were 24.3+, 50+, 10.6, and 78.6+ weeks. For patients with plasma cell myeloma, stable disease was the best response in 17 (63%) patients. The response lasted a median of 11.4 weeks (range, 3.1–46.1 weeks).[74]

Table 2 lists numerous ongoing clinical trials of nivolumab given as a single therapy or in combination with other therapies.

Other anti-PD-1 antibodies

Clinical trials are underway with other anti-PD-1 antibodies (AMP-514, PDR001, REGN2810, BGB-A317, PF-06801591 and AMP-224), but results have not been publicized yet (Table 2).

Pidilizumab (CT-011, MDV9300, previously CT-AcTibody or BAT)

Pidilizumab was originally regarded as an anti-PD-1 antibody. However, Medivation, the company holding the right of pidilizumab, announced via a U.S. Securities and Exchange Commission filing in January 2016 that pidilizumab is not an inhibitor of PD-1.[75] The Food and Drug Administration placed a partial clinical hold on a phase 2 clinical trial of the drug in patients with relapsed or refractory DLBCL, but it lifted the hold in March 2016.[76]

Although the drug showed positive results in patients with follicular lymphoma and DLBCL, its mechanism of action needs to be elucidated.[77, 78]

Anti-PD-L1 antibodies

There are several ongoing clinical trials with anti-PD-L1 antibodies (atezolizumab, durvalumab, avelumab and CA-170) in patients with lymphoid neoplasms, but results have not been published yet (Table 2).

Anti-PD-1/PD-L1 agents in combination with other treatments

Anti-PD-1/PD-L1 antibodies can be combined with other treatments in a myriad of ways. Clinical trials of anti-PD-1/PD-L1 antibodies combined with standard chemotherapy, targeted therapy (i.e., rituximab or BV) or ASCT are plentiful (Table 2). In this review, we will discuss several approaches to combination therapy with anti-PD-1/PD-L1 antibodies.

One of the strategies is to combine an anti-PD-1 antibody with an agent that blocks another co-inhibitory molecules (e.g. CTLA-4 or lymphocyte activation gene 3 [LAG-3]) in T cells. Clinical trials of nivolumab combined with ipilimumab (an anti-CTLA-4 inhibitor) or BMS-986016 (an anti-LAG-3 inhibitor) are underway in patients with hematologic malignancies (Table 2). A similar approach is to combine anti-PD-1/PD-L1 antibodies with urelumab, a cytotoxic T cell- activating drug that binds CD137. The intent of this approach is not to block two different co-inhibitory molecules; rather, it is to simultaneously enhance CTL activity and block inhibitory signals. Preclinical studies have demonstrated that this combination of anti-PD-1/PD-L1 antibodies and urelumab enhances the anti-tumor activity of T cells.[26, 79] A phase 1 clinical trial of nivolumab plus urelumab is now under way in patients with advanced B-cell non-Hodgkin lymphoma.

Combination therapy with anti-PD-1/PD-L1 antibodies can involve modulating the immunosuppressive tumor microenvironment. Epigenetic modifying agents disrupt the immunosuppressive tumor microenvironment by eradicating myeloid-derived suppressor cells or enhancing the effector function of T- and NK cells.[80, 81] Furthermore, treating a human leukemia cell line with a hypomethylating agent has been shown to augment PD-L1 and PD-L2 expression.[82] Therefore, combining epigenetic modifying agents with anti-PD/PD-L1 antibodies is a sensible approach. Indeed, preclinical and clinical studies report promising results.[83, 84] Several clinical trials are under way (Table 2).

In a melanoma mouse model, the tumor-intrinsic, active Wnt/β-catenin pathway induces T cell exclusion in the tumor microenvironment and resistance to anti-PD-L1 antibodies.[85] The kinases FAK and PYK2 augment Wnt/β-catenin pathway, and a preclinical study has shown that combining a FAK/PYK2 dual inhibitor (VS-4718) with an anti-PD-1 monoclonal antibody is more effective than anti-PD-1 therapy alone and extended survival in vivo.[86] Combining VS-4718 and an anti-PD-1 monoclonal antibody also increases the CD8+ T cell: regulatory T cells ratio, suggesting an attractive approach to modulating the tumor microenvironment to enhance the anti-tumor activity of anti-PD-1 antibodies.[87] Patients with plasma cell myeloma could be potential candidates because VS-4718 alone already has been shown to inhibit myeloma cell growth in vitro and in vivo.[88]

It is noteworthy that the administration of AFM13 and an anti-PD1 agent using an autologous PDX mouse model with donor-matched tumors and peripheral blood mononuclear cells from patients with Hodgkin lymphoma showed synergistic anti-tumor effect. AFM13 is a bi-specific, anti-CD30/CD16A, tetravalent chimeric antibody construct that targets CD30-expressing malignancies by recruiting NK cells.[89] Compared to monotherapy with anti-PD1, combined therapy with AFM13 initially enhances the infiltration of macrophages and activated NK cells and later enhances T cells and dendritic cell infiltration.[90] Based on this information, a clinical study of combined AFM13 and pembrolizumab in relapsed/refractory CHL is in preparation.

Chimeric antigen receptor (CAR) T cell therapy has been shown to be effective in patients with lymphoma, and blocking the PD-1 pathway in combination with CAR T cell therapy is an interesting approach.[91] A phase 1 study of durvalumab and anti-CD19 CAR T cell therapy will soon enroll patients with relapsed/refractory DLBCL. Oncolytic viral therapy activates innate immune responses against virally infected tumor cells and enhances adaptive anti-tumor immune responses by in vivo priming against tumor-associated antigens. A melanoma mouse model study of oncolytic vital therapy combined with anti-PD-1 or anti-CTLA-4 demonstrated significant anti-tumor activity, providing a rationale for clinical studies.[92]

Immune-related adverse events

Thanks to the clinical success of nivolumab and pembrolizumab in solid tumors, the list of tumors treatable with anti-PD-1 or anti-PD-L1 antibodies is expanding. Therefore, clinicians would observe more patients with immune related adverse events (irAEs). Overall, grade 3 or 4 irAEs are observed in 7–12% of patients with solid tumors who receive single anti-PD-1 or anti-PD-L1 antibodies.[39, 93] Of note, a predictable pattern of irAEs has been observed in such patients; dermatologic and gastrointestinal toxicities appear early, and hepatic toxicities or endocrinopathies are seen later.[94]

In patients with lymphoid neoplasms, irAEs of any grade appear in 72%–100% of patients. [37, 67, 69, 72–74] Common irAEs include thrombocytopenia, neutropenia, fatigue, infusion reaction, hypothyroidism, rash, diarrhea, nausea, pyrexia, pneumonitis, diarrhea, fatigue, back pain, decrease in platelets, dry skin, and couth.[67–69, 72] Grade 3 or higher irAEs are observed in 11–22% of patients and include interstitial pneumonia, pneumonitis, colitis, gastrointestinal inflammation, increased alanine aminotransferase/aspartate aminotransferase levels, pancreatitis, nephrotic syndrome, fulminant type 1 diabetes mellitus, myelodysplastic syndrome, leukopenia, thrombocytopenia, septic meningitis, pyrexia, infusion reaction, joint swelling, pain, stomatitis, tumor progression, and arrhythmia.[37, 67, 72–74]

The management of irAEs depends on their severity. Patients with grade 1 irAEs may continue PD-1-targeted therapy if symptoms are not observed, but it is recommended to withhold anti-PD-1 therapy from those with grade 2 irAEs and to manage symptoms with oral prednisone (1mg/kg/day) or an equivalent drug. Patients with irAEs of grade 3 or higher should discontinue anti-PD-1 therapy and be treated with intravenous methylprednisolone

(2–4mg/kg/day) or an equivalent drug. Daily monitoring with liver function test is also recommended.[95]

Conclusion

Programmed death-1 (PD-1) is a co-inhibitory molecule and is seen in CD4+ and CD8+ T cells. Upon binding to its ligands, programmed death ligand-1 (PD-L1) and -2 (PD-L2), PD-1 negatively regulates interleukin 2 (IL-2) production and T cell proliferation. PD-L1 expression in normal tissue is limited, but its expression in tumor cells can be induced by extrinsic signals (e.g., IFN-γ). In various lymphoid neoplasms, PD-L1 expression can also be induced by intrinsic signals, including 1) genetic aberrations involving 9p24.1 encompassing $PD-L1$, $PD-L2$ and $JAK2$, 2) latent EBV infection, particularly by LMP1; 3) PD-L1 3′-UTR disruption; and 4) the activated JAK/STAT pathway. Clinical use of PD-1 pathway-blocking agents has successfully treated some lymphoid neoplasms, particularly those with PD-L1 expression induced by intrinsic signals. Currently, combination therapies involving anti-PD-1/PD-L1 agents and conventional chemotherapies, targeted therapies, or other immunotherapies are being studied, and we expect that the resulting data will broaden our understanding of the PD-1 pathway and expand the list of patients who will benefit from PD-1-pathway-blocking agents to include those who suffer from lymphoid neoplasms.

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Highlight points

- **•** Immune checkpoint molecules are deregulated in hematological malignancies through diverse mechanisms
- **•** Checkpoint antagonists have shown encouraging therapeutic effects in patients with relapsed/refractory classical Hodgkin lymphoma, follicular lymphoma and diffuse large B cell lymphoma
- **•** Combined checkpoint antagonists with agents 1. reversing T cell dysfunction, 2. regulating compensatory immune pathway and 3. enhancing tumor antigen are under robust clinical evaluation and shows great promise
- **•** Checkpoint blockade may emerge as a potential agent for consolidation or salvage therapy for both autologous and allogeneic stem cell transplantation

Practical points

• Clinical value of PD-1 signaling pathway dysregulation and regimen selection

- **•** Relationship between monotherapy or combination regimen and potential complications
- **•** Development of clinical algorithm for patient evaluation

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Figure 1.

The two-signal model of T-cell activation. The first signal is a processed antigen presented by the MHC molecule of an APC to TCR of a T cells. This triggers TCR signaling, which is modulated by the antigen-independent co-stimulatory or co-inhibitory signals delivered by APCs. The T cell molecules CD28, PD-1, and CTLA-4 bind to B7-1/B7-2, PD-L1/PD-L2 and B7-1/B7-2, respectively. CD28, when engaged with its ligands, induces cell cycle progression, interleukin-2 production and clonal expansion. In contrast, PD-1 and CTLA-4 induces T cell tolerance when engaged with their respective ligands.

APC, antigen-presenting cells; CTLA-4, cytotoxic T-lymphocyte-associated protein 4; MHC, major histocompatibility complex; TCR, T cell receptor; PD-1, programmed death 1; PD-L1/2, programmed death-ligand 1 or 2, +, activation of the pathway; –, inhibition of the pathway.

Figure 2.

PD-1 and its downstream effect. Upon binding to ligands, PD-1's ITIMs and ITSMs are phosphorylated by Src-family tyrosine kinases. The phosphorylated tyrosine residue subsequently recruits SHP-2 and SHP-1/SHP-2 in ITIM and ITSM, respectively. Activated PD-1 eventually hinders PI3K/Akt and RAS/MEK/ERK pathways, thwarts the function of PKC-θ and ZAP70 phosphorylation and inhibits glycolysis. The net effect is decreased cell cycle progression, IL-2 production, T-cell activation and effector T-cell development and increased apoptosis.

Src, Src-family tyrosine kinases; ITIM, immunoreceptor tyrosine-based inhibitory motif; ITSM, immunoreceptor tyrosine-based switch motif; P in red circle, phosphorylated tyrosine residues; SHP1 and SHP2, Src homology 2 domain-containing phosphatases, PI3K/Akt, Phosphatidylinositol-4,5-bisphosphate 3-kinase; Akt, Protein kinase B; PKC-θ, protein kinase C-theta; RAS/MEK/ERK, RAS/MEK/ERK pathway.

Figure 3.

Images of various lymphoid neoplasms with PD-L1 expression. A–B. Diffuse large B-cell lymphoma, not otherwise specified. Hematoxylin and eosin (A, x400) and PD-L1 stain (B, x400). C–D. Primary mediastinal large B-cell lymphoma. Hematoxylin and eosin (C, x400) and PD-L1 stain (D, x400). E–F. Epstein-Barr virus-positive diffuse large B-cell lymphoma, not otherwise specified. Hematoxylin and eosin (E, x400) and PD-L1 stain (F, x400). G–H. Primary central nervous system lymphoma. Hematoxylin and eosin (G, x400) and PD-L1 stain (H, x400). I–J. Primary testicular lymphoma. Hematoxylin and eosin (I, x400) and PD-L1 stain (J, x400). All PD-L1 stains were performed using SP142 clone (Spring Bioscience, Pleasanton, CA, USA).

Table 1

Available clones for anti-PD-L1 monoclonal antibodies

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Clinical trials involving anti-PD-1 or anti-PD-L1 antibodies Clinical trials involving anti-PD-1 or anti-PD-L1 antibodies

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agonist; Daratumumab, a monoclonal anti-CD38 antibody; HIV, human immunodeficiency virus; R/R, relapsed and refractory; NHL, non-Hodgkin lymphoma; CHL, classical Hodgkin lymphoma; DLBCL, diffuse large B-cell lymphoma; FL, follicular lymphoma; MCL, mantle cell lymphoma; MF/SS, mycosis fungoides/Sezary syndrome; PMBL, primary mediastinal (thymic) large B-cell lymphoma; PCNSL, Ibrutinib, a Bruton's Tyrosine kinase inhibitor; Rituximab, a monoclonal antibody against the protein CD20; ASCT, autologous stem cell transplant; BV (Brentuximab vedotin), an antibody-drug conjugate agonist; Daratumumab, a monoclonal anti-CD38 antibody; HIV, human immunodeficiency virus; R/R, relapsed and refractory; NHL, non-Hodgkin lymphoma; CHL, classical Hodgkin lymphoma; DLBCL Ibrutinib, a Bruton's Tyrosine kinase inhibitor; Rituximab, a monoclonal antibody against the protein CD20; ASCT, autologous stem cell transplant; BV (Brentuximab vedotin), an antibody-drug conjugate diffuse large B-cell lymphoma; FL, follicular lymphoma; MCL, mantle cell lymphoma; MF/SS, mycosis fungoides/Sezary syndrome; PMBL, primary mediastinal (thymic) large B-cell lymphoma; PCNSL primary central nervous system lymphoma; PTCL, peripheral T-cell lymphoma; PCM, plasma cell myeloma; CLL/SLL, chronic lymphocytic leukemia/small lymphocytic lymphoma; MZL, marginal zone primary central nervous system lymphoma; PTCL, peripheral T-cell lymphoma; PCM, plasma cell myeloma; CLL/SLL, chronic lymphocytic leukemia/small lymphocytic lymphoma; MZL, marginal zone (GITR: glucocorticoid-induced tumor necrosis factor receptor); BGB-3111, a Bruton's Tyrosine kinase inhibitor; Obinutuzumab, a humanized anti-CD20 monoclonal antibody; Polatuzumab vedotin, an monoclonal antibody (an immune checkpoint inhibitor); AVD, combination of doxorubicin, vinblastine and Dacarbazine; MEDI-551, an anti-19 monoclonal antibody; GWN323, an anti-GITR antibody (GITR: glucocorticoid-induced tumor necrosis factor receptor); BGB-3111, a Bruton's Tyrosine kinase inhibitor; Obinutuzumab, a humanized anti-CD20 monoclonal antibody; Polatuzumab vedotin, an anti-CD79b antibody-drug conjugate; Tremelimumab, an anti-CTLA-4 antibody; AZD9150, an anti-STAT3 inhibitor; CAR-T, chimeric antigen receptor T-cell therapy; Poly ICLC, a toll-like receptor-3 anti-CD79b antibody-drug conjugate; Tremelimumab, an anti-CTLA-4 antibody; AZD9150, an anti-STAT3 inhibitor; CAR-T, chimeric antigen receptor T-cell therapy; Poly ICLC, a toll-like receptor-3 monoclonal antibody (an immune checkpoint inhibitor); AVD, combination of doxorubicin, vinblastine and Dacarbazine; MEDI-551, an anti-19 monoclonal antibody; GWN323, an anti-GITR antibody antibody targeting CTLA-4; Lirilumab, an anti-KIR monoclonal antibody; RRx-001, a pan-epigenetic anticancer agent; Urelumab, an anti-CD137 monoclonal antibody; BMS-986016, an anti-LAG-3 antibody targeting CTLA-4. Lirilumab, an anti-KIR monoclonal antibody; RRx-001, a pan-epigenetic anticancer agent; Urelumab, an anti-CD137 monoclonal antibody; BMS-986016, an anti-LAG-3 thalidomide; Ubilituximab (TG-1101 or UTX), a novel, chimeric monoclonal antibody targeting a unique epitope on the CD20 antigen; TGR-1202, a PI3K δ inhibitor; Idelalisib, a PI3K δ inhibitor; directed to the protein CD30; lenalidomide, a derivative of thalidomide; R-CHOP; combination of rituximab, cyclophosphamide, doxorubicin, vincristine and prednisone; ipilimumab, a monoclonal thalidomide; Ubilituximab (TG-1101 or UTX), a novel, chimeric monoclonal antibody targeting a unique epitope on the CD20 antigen; TGR-1202, a PISK & inhibitor; Idelalisib, a PISK & inhibitor; directed to the protein CD30; lenalidomide, a derivative of thalidomide; R-CHOP; combination of rituximab, cyclophosphamide, doxorubicin, vincristine and prednisone; ipilimumab, a monoclonal INCB024360 (Epacadostat), an indoleamine 2,3-dioxygenase 1 inhibitor; ACP-196 (Acalabrutinib), a more selective, irreversible Bruton's Tyrosine kinase inhibitor; Pomalidomide, a derivative of INCB024360 (Epacadostat), an indoleamine 2,3-dioxygenase 1 inhibitor; ACP-196 (Acalabrutinib), a more selective, irreversible Bruton's Tyrosine kinase inhibitor; Pomalidomide, a derivative of Dinaciclib (MK-7965), a cyclin-dependent kinase inhibitor; AFM13, a bi-specific, tetravalent chimeric antibody construct against CD30 and CD16A; G100, a potent toll-like receptor-4 agonist; Dinaciclib (MK-7965), a cyclin-dependent kinase inhibitor; AFM13, a bi-specific, tetravalent chimeric antibody construct against CD30 and CD16A; G100, a potent toll-like receptor-4 agonist; lymphoma; SCT, stem cell transplant; CNS, central nervous system. lymphoma; SCT, stem cell transplant; CNS, central nervous system.