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# PD-L1 small-molecule modulators: a new hope in epigenetic-based multidrug cancer therapy?

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**Keywords:** immunotherapy; PD-L1; epigenetics; cancer; therapy escape; multitarget approach.

**Teaser:** This article reviews the latest findings regarding PD-L1 immunotherapy related to epigenetically influenced pathophysiological conditions and the state of the art in drug design of PD-L1-targeting agents and combinatory therapies.

**Programmed death-ligand 1 (PD-L1) is an immune checkpoint protein the overexpression of which results in an inhibitory signal that induces T cell exhaustion responsible for immune escape in tumors. Immunotherapy strategies targeting the PD-L1 pathway have achieved remarkable success in treating various types of cancer. More recently, numerous advances in understanding the complex PD-L1 biology have been made, and the first small-molecule inhibitors have been described in the literature. In this review, we highlight the most promising recent advances in understanding the complex regulation mechanisms focusing on small-molecule modulators, which could be used in rational therapy combinations with other epigenetic chemotherapeutic agents.**

## Introduction

Over the past few years, there has been significant progress in cancer immunotherapy<sup>1</sup> and immune checkpoints are now considered key players in cancer biology because tumor cells are able to evade immune surveillance by distinct immune checkpoint pathways.<sup>2</sup> PD-L1, also known as cluster of differentiation 279 (CD279), acts as an immune checkpoint by negatively regulating T cells via the suppression of the PI3K/AKT and Ras/MAPK/ERK signaling pathways.<sup>3,4</sup> PD-L1 is expressed mainly in peripheral tissues and the tumor microenvironment (TME).<sup>5,6</sup> Activation of the PD-L1 signaling pathway is crucial for the regulation of T cell activation, proliferation, and apoptosis, thus having a direct influence on the cell-mediated immune response.<sup>7</sup> PD-L1 is often highly expressed in tumor-infiltrating lymphocytes, B cells, natural killer (NK) cells, monocytes, and dendritic cells (DCs), and its overexpression results in an inhibitory signal that induces T cell exhaustion,<sup>8</sup> responsible for immune escape in tumors.<sup>9</sup> Thus, PD-L1 acts as a type of tumor cell protector, shielding them from T cell-mediated elimination.<sup>10</sup> Conversely, under physiological conditions, PD-L1 is responsible for immune tolerance, maintenance of immune homeostasis, and protection from excessive autoimmune damage to healthy tissues.<sup>10</sup>

Immunotherapy strategies targeting the PD-L1 pathway have been successfully used to treat numerous types of cancer.<sup>1,11</sup> The PD-L1/PD-1 axis has a fundamental role in escaping the immune system in the advanced stages of Hodgkin's lymphoma,<sup>12</sup> non-small-cell lung cancer (NSCLC),<sup>13</sup> renal cancer, urothelial carcinoma,<sup>14</sup> and melanoma.<sup>15</sup> This implies that immune checkpoint inhibitors (ICIs), capable of blocking the activity of immune checkpoints, could result in the release of the immune brake, reactivation of the immune response of T cells to the tumor cells, and finally, antitumor effects.<sup>16</sup> More than 1000 clinical trials have evaluated anti-PD-1/anti-PD-L1 antibodies, and some of which, such as Keytruda (pembrolizumab), Opdivo (nivolumab), Libtayo (cemiplimab), Tecentriq (atezolizumab), Bavencio (avelumab), and Imfinzi (durvalumab), have been approved by the US Food and Drug Administration (FDA) for more than ten cancer indications.<sup>12</sup>

Following considerable advances in the characterization of PD-L1 biology, numerous small molecules have appeared in both patents and in the literature (reviewed in <sup>17-19</sup>). In principle, small molecules are preferred over antibodies because they have significant advantages, such as stability, membrane permeability, non-immunogenicity, and the possibility of oral administration.<sup>20</sup> Their main drawback is that most compounds follow a limited pharmacophore containing a biphenyl structure. In addition, relatively little is known regarding their pharmacological modulation via small molecules<sup>21</sup>; most small molecules have only been reported as patents, with relatively few data generally available,<sup>19,22</sup> and even fewer compounds having reached clinical trials.<sup>19,22</sup> The most advanced compounds are CA-170 (Phase I completed for solid tumors or lymphomas, no data yet available), GS-4224 or Evixapodlin (Phase I completed, with promising data in patients with solid tumors, Phase II ongoing, NCT04049617) and INCB086550 (Phase I preliminary promising data published, Phase II ongoing in patients with solid tumors) with partially promising data available.<sup>23-25</sup>

Tumors and immune systems in patients are very diverse; blocking of PD-L1 still has slow and low (20–40%) response rates, indirectly controlling cancer growth.<sup>26</sup> Recent published evidence underlined that, for an efficient anti-PD-L1 therapy, an intact

integrated immune cycle needs to be established, because damage in any step of this cycle often leads to therapy failure.<sup>27</sup> This cycle has three crucial steps: (i) cancer-antigen presentation and T cell activation; (ii) trafficking of T cells to tumors; and (iii) T cell-mediated recognition and elimination of tumor cells.<sup>28</sup>

There is growing evidence that an impaired immune cycle can be rescued by other chemotherapeutic agents, especially epigenetic modulators; thus, a rational combination or a hybrid approach of PD-L1 modulation with modulators of epitargets could improve the success of checkpoint blockade therapies in the clinic.<sup>11</sup>

Epigenetic modifications include DNA methylation, histone modification, and RNA modifications. Aberrant epigenetic patterns have been associated with the onset and development of many types of cancer.<sup>29</sup> Numerous epimodulators, such as inhibitors of histone deacetylases (HDACs), DNA methyltransferases (DNMTs), lysine-specific demethylase 1 (LSD1), bromodomains, or enhancer of zeste homolog 2 (EZH2) (Figure 2), have been developed in preclinical and clinical settings based on the growing understanding of the plethora of epigenetic mechanisms.<sup>30–32</sup> Epigenetic patterns are altered not only in cancer cells, but also in immune cells.<sup>33</sup> This dysregulation is likely to be crucial for the immunogenic deficiency of cancerous cells and the subsequent augmentation of more immunosuppressive cells.<sup>34</sup> Epigenetic modifications can activate silent genes, which can be immune checkpoint regulators that either augment the immune response or lead to immune evasion.<sup>35</sup> As outlined for PD-L1, monotherapy outcomes with epidrugs were satisfactory in hematological malignancies, whereas solid tumors often showed unsatisfactory results.

Importantly, there is increasing evidence that epigenetic modulators can influence each step of the immune cycle, from antigen presentation and T cell regulation to the modulation of the immunosuppression of cancerous cells.<sup>36</sup> For example, cytokines can be epigenetically downregulated, leading to lower infiltration rates of T lymphocytes; at the same time, endothelin receptors are overexpressed, resulting in cancer progression and angiogenesis.<sup>27</sup>

Strikingly, there are currently numerous clinical trials involving PD-L1 antibodies associated with epigenetic modulators (see<sup>37</sup> for details). In this review, we critically discuss recent findings to stimulate researchers to develop novel potential therapeutic concepts using PD-L1 inhibitors alone or in combination with epigenetic modulators.

### Epigenetic targets and PD-L1 involvement

To date, numerous epigenetic targets, such as HDACs, DNMTs, LSD1, bromodomains, or EZH2 have been linked with the immunomodulation of PD-L1 (Figure 2).

#### HDACs

HDACs are divided into classes I, IIa, IIb, III, and IV, and act mainly via the removal of acetyl groups from the *N*-acetyl lysine amino acid residues of histones,<sup>38</sup> modulating gene expression and regulating PD-L1 expression.<sup>39</sup> Numerous HDAC inhibitors (HDACis) have been developed and are well reviewed elsewhere.<sup>40</sup> Apart from a direct cytotoxicity in cancer, HDACis were demonstrated to change immunogenicity and to enhance antitumoral immune responses by decreasing myeloid-derived suppressor cells (MDSCs) while simultaneously increasing the expression of the co-stimulatory molecules MHC-I/II, CD40, CD80, and CD86.<sup>41,42</sup>

Numerous HDACis, such as panobinostat (pan-HDACi), entinostat (class I HDACi), mocetinostat (HDAC 1,2,3 and 11 inhibitor) and tucidinostat, the first approved (in China) oral class I-selective HDACi, have been shown to increase the expression of PD-L1, mainly in melanoma models,<sup>40,43</sup> while showing a preference for class I selective agents. Most clinical data relating to the successful combination of PD-L1 inhibition and an HDACi relate to the pairing of entinostat and pembrolizumab (NCT02697630), resulting in a durable response in metastatic uveal melanoma.<sup>44</sup> Other combinations of entinostat with the bifunctional (TGF- $\beta$ /PD-L1) fusion protein M7824 (NCT04708470), avelumab (NCT02915523), or nivolumab (NCT03838042) remain under evaluation.

In a clinical setting, Munster *et al.* investigated a triple combination of tamoxifen with the pan-HDACis vorinostat and pembrolizumab for the treatment of hormone-resistant breast cancer, with promising outcomes in some patients (NCT02395627),<sup>45</sup> whereas, for another triple therapy approach with entinostat, nivolumab, and ipilimumab (NCT02453620), no data are yet available. Promising first data are available for a Phase Ib study with vorinostat and pembrolizumab in patients with metastatic prostate carcinoma (NCT02619253).<sup>46</sup> There have been several studies colorectal cancer (CRC), but the results have yet to be reported.<sup>37,47</sup>

The selective inhibition of HDAC3 with RGFP966 increased the expression of PD-L1 by increasing histone acetylation on the PD-L1 promoter, which is the probable cause of resistance to RGFP966. Indeed, subsequent co-treatment with the anti-PD-L1 antibody BE101 enhanced the efficacy of RGFP966 in B cell lymphoma cells as well as in a mouse model.<sup>48</sup> By contrast, in pancreatic cancer cells, the same HDAC3i inhibitor reduced PD-L1 expression by modulating signal transducer and activator of transcription 3 (STAT3), indicating that HDAC3 inhibition could enhance immunotherapy.<sup>49</sup>

Recently, it was demonstrated that HDAC3 overexpression via the constitutive photomorphogenic 1 (COP1)/c-Jun axis inhibited PD-L1 expression in cisplatin-resistant NSCLC cells.<sup>50</sup> Taken together, HDAC3 appears to be able to up or downregulate histone acetylation of the PD-L1 promoter region; thus, further studies are necessary to determine whether selective HDAC3i is a suitable strategy for multidrug immunotherapy.

Similarly, HDAC6 can cause PD-L1 overexpression by modulating the STAT3 pathway, and HDAC6i with tubastatin A and nexturastat A impaired tumor progression in a melanoma mouse model.<sup>51</sup> A subsequent study by another research team provided insights into the mechanism of nexturastat A, which is able to increase the efficacy of anti-PD-L1 antibodies (CD279 and BE0146) by activation of immune surveillance cells and tumor-associated macrophages (TAMS), and not directly over T cells.<sup>52</sup> MPT0G612, another selective inhibitor of HDAC6, blocked IFN $\gamma$ -mediated upregulation of PD-L1 in colorectal carcinoma cells, resulting in apoptosis via autophagy inhibition.<sup>53</sup>

Despite some available clinical data, our understanding of the effects of the combination of HDACis with immunotherapy, especially anti-PD-L1 therapy, remains limited. Further studies are necessary to understand better the complex biology behind these effects, as well as to develop better treatment options. Nevertheless, targeting HDACs appears to be a promising and effective approach to improving the response of the immune system to cancer cells.

*DNMTs*

DNMTs are important epigenetic enzymes, proven to be involved in the onset and development of numerous cancers.<sup>54</sup> To date, various DNMT inhibitors (DNMTis) have either been approved or are under clinical development. Interestingly, these inhibitors can upregulate MHC class I and II in many cancers,<sup>55</sup> such as decitabine in lymphoma. Furthermore, treatment with DNMTis resulted in the expression of several cancer testis antigens, which are cancer-associated antigens directly expressed on these malignant cells, of skin, lung, or colon cancers.<sup>11,56</sup> Thus, DNA methylation is likely a key player in the regulation of their expression. DNMTs also appears to have a crucial role in the maturation and differentiation of DCs because hypomethylated DNA enhancer regions are related to binding domains for transcription factors known to influence the DC lineage.<sup>57</sup>

Patients with resistance to PD-L1/PD-1 antibody treatment have low levels of activated immune cells; thus, activation of immune cells by DNMTis might boost the activity of PD-L1/PD-1 treatment. Indeed, combinations of DNMTis and anti-PD-L1 immunotherapeutic agents are under clinical investigation. Azacitidine is being tested in combination with durvalumab against metastatic ER<sup>+</sup>/HER2<sup>-</sup> breast cancer (NCT02811497), whereas decitabine is being evaluated in co-treatment with pembrolizumab in locally advanced HER2<sup>-</sup> breast cancer (NCT02957968). The latter combination is also being tested on NSCLCs and esophageal carcinomas (NCT03233724).

Recently, the HDAC3i RFGP966 was reported to downregulate DNMT1 expression, resulting in enhanced PD-L1 levels.<sup>48</sup> The proposed mode of action of RFGP966 is a fascinating example of a complex epigenetically driven interplay of three targets: the cascade begins with direct HDAC3 inhibition and subsequent DNMT1 degradation, resulting in increased PD-L1 expression, which renders the tumor more susceptible to PD-1/PD-L1 blockade therapy. It is likely that HDAC3 inhibition does not directly influence DNA methylation of the PD-L1 promoter but instead triggers an IFN response via DNMT1 degradation, resulting in activated PD-L1 expression. However, further studies are necessary to confirm this mechanism.<sup>58</sup> Thus, it is of interest to study whether the acetylation levels of regulatory sequences for PD-L1 also correlate with their methylation.

However, the combination DNMT/PD-L1 inhibition is still in its infancy; thus, more studies are required to understand the potential of this novel combined therapeutic approach.

*LSD1*

LSD1, also known as KDMA1, is an essential epigenetic target mainly studied in the context of cancer. It is a histone demethylase able to modulate the methylation patterns of H3K4 and H3K9. Several inhibitors are currently in clinical trials for numerous forms of cancer, such as acute myeloid leukemia and solid cancers (e.g., breast or lung cancer).<sup>59</sup> Sheng *et al.* reported that PD-L1 inhibition, together with LSD1 ablation, might be a successful therapeutic strategy. Indeed, they were able to show that LSD1 depletion led to tumor immunogenicity and T cell infiltration in poorly immunogenic tumors, such as melanoma, restoring sensitivity to anti-PD-1 therapy. Despite these encouraging results, no pharmacological LSD1 inhibitor has received approval as yet.<sup>60</sup>

Moreover, LSD1 appears to have a crucial role in stemness maintenance and chemoresistance in breast cancer.<sup>61</sup> It is capable of transforming TAMs into M1-like macrophages, probably via the PD-L1 axis, in triple-negative breast cancer<sup>62</sup>; however, the involvement of PD-L1 remains rather elusive.

Interestingly, SP-2577, a novel reversible LSD1 inhibitor, enhanced antitumor immunity and T cell infiltration in an ovarian cancer model of the SWItch/Sucose-NonFermentable (SWI/SNF) mutated complex. This compound activated IFN-dependent antitumor immunity and led to increased expression of PD-L1. A subsequent co-treatment with an anti-PD1 antibody resulted in a significantly enhanced immune response and decreased cancer cell growth.<sup>63</sup>

Besides the relevance in cancer, LSD1 has been demonstrated to be recruited to the PD-L1 gene locus via B lymphocyte-induced maturation protein 1 (Blmp-1) in viral infections. Thus, the removal of H3K4 methylation patterns via LSD1 inhibited expression of PD-L1.<sup>64</sup> Thus, a combination of LSD1 inhibitors together with an anti-PD-1 therapy is of high interest not only in regard of cancer, but also other immune-system-related diseases to increase therapy success and decrease resistance.

*Bromodomains*

Bromodomain and extraterminal (BET) motif inhibition is now considered a potential strategy in epigenetic-driven cancer therapy, but to date, no drugs have been approved despite intensive research.<sup>65</sup> BET proteins can control PD-1 expression in T cells as well as PD-L1 expression in cancer cell models. For example, BET inhibitors, such as JQ1, were shown to differentiate effector T cells into effector memory cells via suppression of the transcription factor BATF in a leukemia cell model.<sup>66</sup> PD-L1 is a target of JQ1 in neuroblastoma,<sup>67</sup> underlining the importance of the connection between PD-L1 and BET. In addition, in lung cancer, BET inhibition by JQ1 disrupted regulatory T cells (Tregs), and a concomitant PD-1 antibody treatment resulted in an improved antitumoral response in a mouse model.<sup>68</sup>

Moreover, promising results were recently published in triple-negative breast cancer,<sup>69</sup> highlighting targeting BET as a promising approach to overcome immune resistance and to improve antitumor therapy responses via reduction of the PD-1/PD-L1 axis. Based on these findings, several clinical trials are evaluating BET inhibitors other than JQ1 (i.e., SF1126, RO6870810, or BMS-986158) in combination with anti-PD-1 or anti-PD-L1 antibodies in patients with different cancers (NCT03059147, NCT03292172, and NCT02419417), further underlining PD-L1 inhibition with BET inhibitors as a valuable therapeutic approach.

*EZH2*

EZH2 is a well-known H3K27 histone methyltransferase, the overexpression of which is associated with numerous types of cancer. Various compounds have been studied as EZH2 inhibitors and, recently, tazemetostat entered the market.<sup>70</sup> Numerous clinical trials are ongoing, mainly limited to hematological malignancies.<sup>71</sup>

The role of EZH2 in immunology is not fully clear, because some studies suggest a positive influence on the immune response to fight cancer, whereas others suggest suppressed antitumor immunity. In 2019, two EZH2 inhibitors, MC4040 and MC4041, were

reported as being able to decrease proliferation in glioblastoma (GBM) cells and also to impair the GBM proinflammatory phenotype, with a significant decrease in TGF $\beta$ , TNF $\alpha$ , and IL6, with a concomitant increase in the anti-inflammatory cytokine IL10.<sup>72</sup>

Peng *et al.* reported that treatment with the EZH2 inhibitor GSK126 reactivated Th1-type chemokines, such as CXCL9 and CXCL10, thus allowing trafficking of effector T cells into the TME.<sup>73</sup> By contrast, EZH2 inhibition with GSK126 promoted the development of MDSCs, which are known to suppress an effective antitumor immune response.<sup>74</sup> EZH2 appears to have a crucial role in the differentiation of peripheral Tregs, which are suspected to influence the efficacy of PD-(L)1 immunotherapy.<sup>75</sup> Although EZH2 appears to be a double-edged sword in immunotherapy, the first clinical trial of a combination of the EZH2 inhibitor tazemetostat with pembrolizumab is underway in urothelial carcinoma.<sup>76</sup> However, more studies are necessary to understand the potential converging pathways of EZH2 and PD-L1 before proceeding with eventual co-treatment strategies.

#### **PD-(L)1 small-molecule inhibition: a new frontier for cancer treatment?**

As outlined above, PD-(L)1-specific antibodies have reached the market; however, despite remarkable positive outcomes in some patients, antibodies have well-known disadvantages when used as therapeutic agents, such as high costs, poor bioavailability, and above all, immune-related side effects<sup>77</sup>; thus, small molecules acting against the PD-1/PD-L1 pathway are urgently needed. Initially, researchers focused on small molecules able to target PD-(L)1, including peptides, peptidomimetics, or even macrocyclic peptides mainly developed by Aurigene or Bristol-Myers Squibb, often only described in patents with little or no research data available.<sup>19</sup> These early PD-(L)1 inhibitors are reviewed elsewhere.<sup>17–19,22,78,79</sup> Interestingly, despite the good validation of PD-(L)1 antibodies *in vitro* and *in vivo*,<sup>22,80</sup> the small molecules, even though very potent biochemically speaking, do not exhibit anticancer activity, or no data are available in this regard.<sup>19,22,81</sup>

Bristol-Myers Squibb, who pioneered the field of PD-(L)1 inhibition, also disclosed in 2015 the first nonpeptidic small molecules able to disrupt the PD-1/PD-L1 complex, exhibiting IC<sub>50</sub> values of 1–300 nM<sup>82,83</sup>; BMS-202 and BMS-1018 with an *m*-(phenyloxymethyl)biphenyl scaffold have been prototypes for numerous subsequent molecules based on this pharmacophore,<sup>22</sup> which was described in detail showing the protein–protein interactions (PPIs) of the human PD-1/PD-L1 complex in detail.<sup>83</sup>

From these crystallographic studies, researchers understood that the binding surface of PD-(L)1 is relatively flat and hydrophobic, making the rational design of small molecules acting as PD-(L)1 inhibitors difficult; however, there is room for improvement because the numerous crystal structures of PD-(L)1 available alone or bound to various monoclonal antibodies<sup>84</sup> can provide inspiration for how to design and improve PD-L1 inhibitors by studying hot-spot residues in the aforementioned crystals in a dynamic environment. Figure 3 shows a schematic of the pharmacophore model as well as the key amino acids interacting with PD-L1 small molecules.

Most current PD-(L)1 modulators follow the precise pharmacophore model of compound 1: the biphenyl group appears to be vital and is the main driver of PD-L1 binding.<sup>85</sup> The ether linker does not appear to be crucial,<sup>86</sup> whereas a flexible central aromatic core connected, mainly in the *meta* or *para* position, to a polar tail appears to be essential for interactions with PD-L1. This polar tail often comprises an aminoethanol or a (substituted) ethylenediamine chain. Studies shed light on the mechanism of action of BMS-202 analogs because this molecule can lead to the formation of PD-L1 protein homodimers, hampering PD-L1 binding to its receptor PD-1 and, thus, blocking downstream signaling.<sup>87,88</sup> This mechanism is also likely to be valid for other related derivatives, reviewed elsewhere.<sup>17–19,22,78,79</sup> We highlight the most recent and/or promising small molecules in Table 1.

#### **Concluding remarks and perspective**

Chemotherapeutic agents have been used for decades as cancer therapy; however, the problems of relapse and resistance remain unsolved. The field of ICIs is rapidly evolving, and targeting PD-(L)1 has been proven to be particularly successful, as indicated by the FDA approval of several monoclonal antibodies<sup>12</sup>; however, the side-effects of antibody therapy are well known in terms of a low response because of intrinsic or acquired resistance, but also adverse immune effects limiting the use of these agents.<sup>93</sup> Safer treatment options are urgently needed, and research is focusing on peptidic and nonpeptidic small molecules acting on PD-(L)1.<sup>94</sup> Such molecules have numerous advantages over monoclonal antibodies, such as stability, membrane permeability, no immunogenicity, and last, but not least, oral administration. However, the development of these small-molecule immune checkpoint modulators is still in its infancy,<sup>95</sup> because no such inhibitor of the PD-(L)1 pathway has yet been approved, although the first promising examples have arrived in the clinic, such as CA-170, GS-4224, and INCB086550, with some promising data available.<sup>23–25</sup>

PPIs, such as PD-1/PD-L1, are challenging to target, as also seen for other PPIs, such as IL-2/IL-2R, p53/MDM2, BCL-2/BAX, GP120/CCR5, and Hsp90/Cdc37.<sup>84</sup> Few of these other targets have been successfully approached by small molecules, underlining the difficulties of the drug development process in this regard. As stated by Deng *et al.*, the discovery and development of most PD-L1 small molecules so far has often relied only on protein-based assays, such as homogeneous time-resolved fluorescence, without a clear translation into a more complex biological system, such as cells or mice; thus, their biochemical activity only might not be sufficient.<sup>19</sup> Previously, drug design has been based on a somehow static protein–ligand complex, but PPIs are highly dynamic and have to be studied as flexible plastic targets. The ample crystal structures of PD-(L)1 available, either alone or bound to various monoclonal antibodies were discussed in a recent perspective,<sup>84</sup> providing inspiration for how to design and improve PD-L1 inhibitors by studying hot-spot residues in the aforementioned crystals in a dynamic environment. For these approaches, advanced computational and biophysical techniques are required. These facts underline why small molecules against PD-L1 are hard to develop, even for big pharma.

Today, most small molecule PD-(L)1 modulators have a biphenyl ether-based moiety,<sup>17,19,79</sup> which appears to be vital for most of the highly active small molecules developed to date. Given that the crystal structures of numerous PD-L1 inhibitors are now available, it would be interesting to determine whether and how this moiety could be replaced to improve activity and pharmacokinetic/pharmacodynamic properties.<sup>22</sup>

Despite being nanomolar in biochemical assays, some PD-L1 inhibitors have not yet been validated in mice<sup>22</sup>; thus, they cannot be thoroughly evaluated. Therefore, the development of new structurally diverse PD-L1 inhibitors is of interest not only as improved

treatment options, but also as tools to understand better the complex biology behind them. Recently, compound 21 was described as a selective chemical probe<sup>92</sup> that will undoubtedly boost our understanding of the complex biology behind PD-L1 interactions.

Reprogramming of tumor cells via epigenetic mechanisms is crucial for escape from immune surveillance.<sup>11</sup> Over the past 30 years, epigenetic modulators have been under active development, resulting in valid therapeutic options<sup>96</sup>; however, the problems of relapse and resistance remain<sup>12</sup>; thus, urgent novel treatment options in the form of combination therapies are necessary.

As outlined above, the various steps in the immune cycle are regulated by epigenetic mechanisms, which can be influenced by epigenetic modulators. Some epidrugs have been demonstrated to improve antigen presentation, T cell trafficking, and migration, ultimately breaking the immunosuppressive state.<sup>36</sup>

As an immune checkpoint in combination with epigenetic therapeutic agents, PD-L1 is able to improve immune recognition and tumor growth, leading to increased response rates in preclinical and clinical studies.<sup>36</sup> Several combination approaches are currently under evaluation in numerous cancers, including colon, prostate, breast, and melanoma. However, these studies are mainly clinical, and the output is often unknown or not (yet) published. Combinations of epimodulators and antibodies against PD-L1 are still in their infancy and few results are available; nevertheless, this approach shows considerable therapeutic potential. However, we believe that research efforts also need to be directed toward evaluation of this combination approach by studying the crossroads of the potentially convergent molecular pathways so that combinations of PD-L1 antibodies and epigenetic modulators can be tested in a clinical setting, supported by sound scientific understanding.

Researchers are beginning to understand the interactions of PD-(L)1 with epigenetic enzymes. RFGP966 is a fascinating example of a complex interplay that results in direct HDAC3 inhibition, subsequent DNMT1 degradation, and increased PD-L1 expression, making the tumor more susceptible to PD-1/PD-L1 blockade therapy<sup>48</sup>; however, further studies are necessary to confirm the indirect mechanisms of action of this compound.<sup>58</sup> To sum up, although many questions remain, our increased understanding over the past few years underlines the fact that the combination therapy of epigenetic agents with immunomodulators is a rapidly evolving topic. In the years to come, we can expect a rapidly expanding knowledge and interest in the field of immune checkpoint therapy and epigenetic agents, from not only a biological, but also a therapeutic viewpoint.

#### Authors' contributions

All the authors contributed to the composition and revision of this article.

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#### Declaration of interests

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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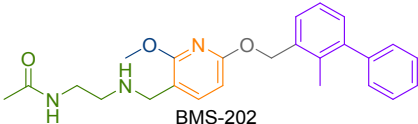
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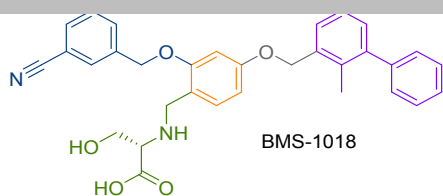
**Figure 1.** Chemical structures and data availability of programmed death-ligand 1 (PD-L1) inhibitors CA-170, GS-4224, and INCB086550 in clinical trials.

**Figure 2.** Epigenetic modulators in combination with programmed death-ligand 1 (PD-L1) antibodies: histone deacetylases (HDACs), DNA methyltransferases (DNMTs), lysine-specific demethylase 1 (LSD1), bromodomains, or enhancer of zeste homolog 2 (EZH2) small-molecule inhibitors currently used in (pre)clinical settings with various PD-(L)1 antibodies. Abbreviation: NSCLC, nonsmall cell lung cancer.

**Figure 3.** Programmed death-ligand 1 (PD-L1) small-molecule inhibitor pharmacophore model with a schematic of the main interactions with the amino acids of the binding site. Amino acids from chain A are depicted in red, whereas those of chain B are in blue.

**Table 1. Summary of the most promising recent PD-L1 small-molecule modulators**

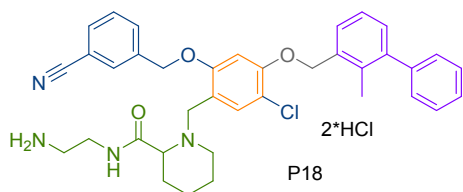
| PD-L1 inhibitor                                                                                | IC <sub>50</sub> | Biological activity                                                                                                                            | Refs           |
|------------------------------------------------------------------------------------------------|------------------|------------------------------------------------------------------------------------------------------------------------------------------------|----------------|
| <br>BMS-202 | 18 nM            | Parent compound for most PD-L1 inhibitors; studied in numerous cellular and mouse models with various success, no transformation to the clinic | 17–19,22,78,79 |



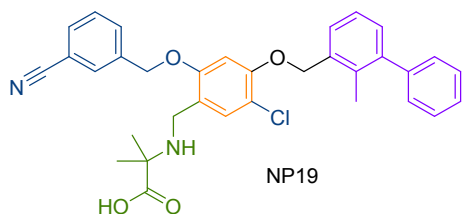
9.1 nM

Active B16-F10 melanoma mouse model

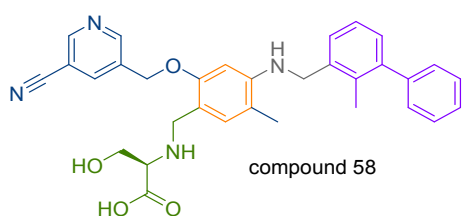
89,90



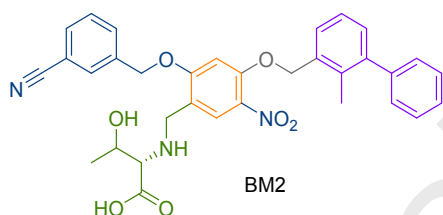
12.9 nM

Active in B16-F10 melanoma mouse model; positively evaluated in H22 model of liver cancer <sup>89,90</sup>

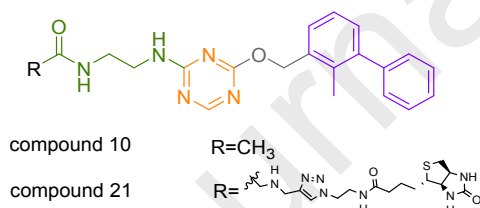
12.0 nM

Active in humanized MC38 colon cancer mouse model, with similar potency to BMS-202 but with less unspecific toxicity <sup>91</sup>

2.7 nM

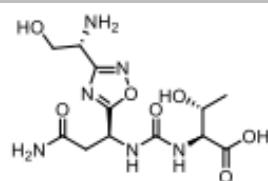
Active in Lewis lung carcinoma *in vitro* and in allograft mouse model, being more potent than BMS-1018 <sup>81</sup>

115 nM for 10 and ND for 21

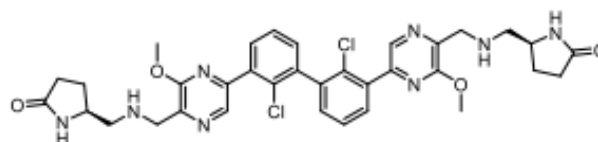
Compound 10 induces apoptosis more strongly than BMS-202 in PC9 or HCC827 lung cancer, despite being biochemically less potent. Compound 21 is first biotinylated probe available to further study its binding capability directly to PD-L1 on cell membranes <sup>92</sup>

### Highlights

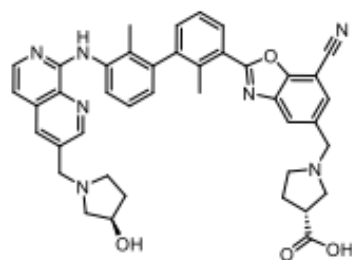
- Short digest on the latest updates regarding immunotherapy with a focus on PD-L1
- Summary of the recent advances in the field of small molecule PD-L1 modulators
- Overview of the growing interest in rational therapy combinations of PD-L1 modulators with epi-drugs



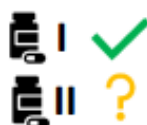
CA-170  
IC<sub>50</sub> 5-10 nM



GS-4224  
IC<sub>50</sub> not yet available



INCB086550  
IC<sub>50</sub> <10 nM



#### Key



Clinical trial Phase I



Clinical trial Phase II



Results available



Unknown/no results available yet

