PD-L1 expression levels on tumor cells affect their immunosuppressive activity

YANG ZHENG1-3*, YOU-CHEN FANG1,4* and JING LI1

1Chinese Academy of Sciences Key Laboratory of Receptor Research, National Center for Drug Screening, Shanghai Institute of Materia Medica, Chinese Academy of Sciences, Shanghai 201203; 2School of Life Science and Technology, ShanghaiTech University, Shanghai 201210; 3School of Pharmacy, University of Chinese Academy of Sciences, Beijing 100049; 4School of Life Sciences, Shanghai University, Shanghai 200444, P.R. China

Received August 24, 2018; Accepted August 6, 2019

DOI: 10.3892/ol.2019.10903

Abstract. Programmed cell death 1 (PD-1) is an immune-checkpoint receptor which is primarily expressed on T cells, monocytes, natural killer cells and macrophages. Programmed death-ligand 1 (PD-L1) is the primary ligand of PD-1 and is constitutively expressed on antigen presenting cells, mesenchymal stem cells and bone marrow-derived mast cells. In addition, PD-L1 is also expressed on a wide range of tumor cells, including lung cancer, breast cancer and melanoma. PD-1 and PD-L1 are important members of the immunoglobulin super-family and participate in immune regulation. In the present study, the immune-suppressive effects of a number of tumor cell lines were determined. The breast tumor cell lines MCF-7 and MDA-MB-231 displayed the largest inhibitory effects on T-cell activation and cytokine secretion in a co-culture system. The HepG2, A549 and A375 cells displayed limited inhibitory effects. MCF-7 and MDA-MB-231 cells expressed the highest level of PD-L1 among the cells used, which may explain their higher immunosuppressive effects. Compound A0-L, a small molecule inhibitor of the PD-1/PD-L1 interaction, restored T cell functions. Additionally, it was demonstrated that the tumor cells with higher levels of PD-L1 expression suppressed signaling pathways involved in T-cell activation, such as the T-cell receptor-zeta chain of T cell receptor associated protein kinase ZAP70-RAS-GTPase-extracellular-signal-regulated kinases and CD28-PI3K-Akt serine/threonine kinases pathways. These findings suggest that tumor cells with higher expression levels of PD-L1 may exhibit higher immuno-suppressive activity, and that drugs targeting the PD-1/PD-L1 interaction may have improved therapeutic effects on tumors expressing higher levels of PD-L1.

Introduction

Tumors utilize a variety of mechanisms to impair the functionality of tumor-specific immune cells, T cells, macrophages and other cells associated with the immune response (1,2). These mechanisms include the expression of ligands which bind to inhibitory receptors expressed on T cells and suppressing the function of T cell stimulatory receptors, such as T-cell receptor (TCR)/CD3 and CD28 (3,4). In general, T cells are activated by the interaction of the TCR/CD3 complex with an antigen and co-activation of CD28 (5). Co-stimulation of the TCR with CD28 and an antigen promotes the initial phosphorylation events of signal transduction from the TCR and enhances immune support functions (6). In addition to the foremost activation pathways, a number of immune checkpoints have been discovered to regulate the immune system. These pathways are crucial for self-tolerance and innate immunity and prevent the immune system from attacking cells indiscriminately (2). Immune checkpoints consist of stimulatory checkpoint molecules and inhibitory checkpoint molecules (7,8). Inhibitory checkpoint molecules have been considered important targets for cancer immunotherapy (9). Currently, several checkpoint inhibitors which block cytotoxic T-lymphocyte associated protein 4 (CTLA4), programmed cell death-1 (PD-1) and programmed death ligand-1 (PD-L1) have been approved for clinical use (10).

The immune system regulates tumor biology, and, depending on the tumor, can either support or inhibit tumor development, growth, invasion and metastasis (11,12). Certain tumors may evade immune detection through recruitment of immunosuppressive leukocytes, which create a microenvironment that blocks the antitumor immune response. Several mechanisms, including defects in antigen-presenting cells, negative immune regulation by suppressive cells and defective antitumor T cells have been hypothesized and demonstrated to explain evasion or tolerance of the immune response in
different types of cancer (11). Jurkat cells are an immortalized line of human T lymphocyte cells that have been used to study acute T-cell leukemia and T-cell signaling (13). Jurkat cells have been used in a diverse array of molecular investigations, some of which underpin our current understanding of multiple signaling pathways (13). Evidence suggests that CD3/CD28-costimulated Jurkat T cells and co-engagement of TCR/CD3 and CD28 results in interleukin (IL)-2 production and activation of extracellular signal regulated kinase (ERK)/c-Jun N-terminal kinase and NF-κB inhibitor β kinase, which is frequently used as a functional readout of activation of Jurkat cells (14).

As an immunosuppressive molecule receptor, PD-1 can inhibit the activation of T lymphocytes and play an important role in immune escape. PD-1 belongs to the CD28/CTLA-4 family of molecules, and it negatively regulates PD-1 signaling. When two PD-L1 or PD-L2 ligands are expressed, PD-1 is downregulated (17,23). PD-1/PD-L1 interaction activates a signal which inhibits TCR-mediated T-cell activation and proliferation, suppresses secretion of cytokines, such as interferon-γ (IFN-γ) and interleukin-2, and promotes cytotoxic T-cell apoptosis and regulatory T-cell differentiation (24,25). A number of pathways involved in T-cell activation, including major histocompatibility complex (MHC)-TCR-ZAP70-RAS-GTPase (RAS)-ERK and CD80-CD28-PI3K protein kinase B (Akt) pathways, have been reported to be regulated by PD-1/PD-L1 interaction.

Although several PD-1/PD-L1 inhibitors have been approved for cancer therapy, the effectiveness of these inhibitors appears to be tumor specific (26). Therefore, the aim of the present study was to determine whether the expression levels of PD-L1 on tumor cells affected its immunosuppressive activity and thus, the therapeutic effects of PD-1/PD-L1 inhibitors. The immuno-suppressive effects of a number of tumor cell lines were assessed. It was demonstrated that MCF-7 and MDA-MB-231 cells expressed the highest levels of PD-L1 and also displayed the highest degree of immune suppression. Additionally, the tumor cells with increased levels of PD-L1 expression exhibited suppression of the pathways involved in T-cell activation, including TCR-ZAP70-RAS-ERK and CD80-CD28-PI3K-AKT pathways.

Materials and methods

Plasmid and cell lines. A PGL3-nuclear factor of activated T-cells (NFAT)-TA-Luciferase plasmid containing the full-length luciferase gene under the control of an NFAT-driven promoter, was used in the present study. NFAT is a nuclear factor of activated T cells that synergizes with activator protein 1 transcription factors at composite sites that are located in the promoters and enhancers of a number of cytokine genes. This indicates that the NFAT promoter an important factor in the immune response (27). The PGL3-NFAT-TA-Luciferase plasmid was generously provided by Dr Jia Li (Shanghai Institute of Materia Medica Chinese Academy of Science). The human embryonic kidney cell line 293, human breast cancer cell line MCF-7, human melanoma cell line A375, human cervical cancer cell line HeLa and human liver cancer cell line HepG2 were obtained from the American Type Culture Collection (ATCC) and maintained in DMEM, supplemented with 10% FBS. The human breast cancer cell line MDA-MB-231, was obtained from ATCC and maintained in Leibovitz’s L-15 medium, supplemented with 10% FBS. The human lung cancer cell line A549, was obtained from ATCC and maintained in DMEM, supplemented with 10% FBS, 1% Glutamax™. The human T lymphocyte cell line Jurkat, was obtained from ATCC and maintained in RPMI-1640 medium, supplemented with 10% FBS, 1% Glutamax™ and 0.1% 2-mercaptoethanol. Human peripheral blood mononuclear cells (PBMCs) were isolated by density gradient centrifugation using a Ficoll-Paque solution (GE Healthcare) from heparinized peripheral blood samples.

Reagents. DMEM, RPMI-1640 medium, Leibovitz 15 medium, FBS, Glutamax™ and 2-mercaptoethanol were purchased from Gibco; Thermo Fisher Scientific, Inc. Penicillin-streptomycin was purchased from Sigma-Aldrich (Merck KGaA). FuGENE® HD Transfection Reagent was purchased from Promega Corporation and TRIZol® reagent was purchased from Invitrogen (Thermo Fisher Scientific, Inc). Anti-CD3 and anti-CD28 were purchased from BD Biosciences. Allophycocyanin-conjugated anti-human-CD274/PD-L1 antibody were purchased from eBioscience (Thermo Fisher Scientific, Inc). Rabbit anti-Erk1/2, rabbit anti-phospho-Erk1/2, rabbit anti-AKT, rabbit anti-GAPDH and horseradish peroxidase-conjugated Goat anti-rabbit immunoglobulin G were purchased from Cell Signaling Technology, Inc. Human IFN-γ ELISA kits and human IL-2 ELISA kits were purchased from Cisbio (PerkinElmer, Inc.). A0-L (a PD-1 inhibitor; patent no. WO 2015/034820 A1; molecular weight, 475.58) was synthesized by Dr Wei Lv of East China Normal University. The company name and catalog number for ELISA kits and all antibodies are listed in Table SI.

Preparation of conditioned medium. A total of 2x10⁵ 293, MCF-7, A375, A549, HeLa or HepG2 cells were cultured per well in 6-well culture plates with DMEM, supplemented with 10% FBS at 37°C for 24 h. In the A549 cells, 1% Glutamax™ was added to the culture medium. MDA-MB-231 (2x10⁵ cells/well) were cultured in 6-well cell culture plates with Leibovitz’s L-15 Medium, supplemented with 10% FBS and 1% Glutamax™ at 37°C for 24 hours. The culture media was collected and centrifuged at 12,000 x g for 10 min using a pre-chilled centrifuge set to 4°C. The supernatant was collected and termed ‘conditioned medium’.
Co-culture of tumor cells with Jurkat cells. Jurkat cells were transfected with 3.3 µg PGL3-NFAT-TA-Luciferase using FuGENE® HD Transfection Reagent. After 16 h, 2x10^5 MCF-7, MDA-MB-231, A549, A375, HeLa or HepG2 cells were seeded into wells with their respective growth medium, and 2x10^4 Jurkat cells transfected with PGL3-NFAT-TA-Luciferase were added to the wells. The conditioned media was collected from the cultures after 24 h. The conditioned media was added to 2x10^4 Jurkat cells transfected with PGL3-NFAT-TA-Luciferase. After 30 min, anti-CD3 (1 µg/ml) and anti-CD28 (1 µg/ml) were added to the culture systems: Tumor cells; Jurkat cells co-cultured with tumor cells in normal media; Jurkat cells alone in conditioned media. After 24 h of co-culture, luciferase activities were measured using the Luciferase system (Promega Corporation) and EnVision multiplate reader (PerkinElmer, Inc.) according to the manufacturer's protocol.

Isolation of human PBMCs. PBMCs derived from healthy volunteers were provided by the Shanghai Blood Center. PBMCs were isolated using a Ficoll-Paque gradient. To separate PBMCs, 20 ml Ficoll was placed in a 50 ml conical centrifuge tube and an equal volume of whole blood diluted 1:1 with PBS was layered on top. The 50-ml tubes were centrifuged at 2,000 x g for 30 min at room temperature with a low acceleration speed. The PBMCs at the interface between the Ficoll and the plasma were gently collected by aspiration using a Pasteur pipette and placed in a 15 ml conical tube. Subsequently, the PBMCs were washed twice with 10 ml PBS and centrifuged at 2,000 x g for 5 min at 4°C (28). The PBMCs at the interface between the Ficoll and the plasma were gently collected by aspiration using a Pasteur pipette and placed in a 15 ml conical tube. Subsequently, the PBMCs were washed twice with 10 ml PBS and centrifuged at 2,000 x g for 5 min at 4°C (28). The PBMCs were cultured for 6 h in T25 flasks in complete RPMI-1640 media (fresh RPMI-1640 medium supplemented with 10% FBS, 2 mM L-glutamine, 100 IU/ml penicillin and 100 µg/ml streptomycin and 0.1% 2-mercaptoethanol) at 37°C in a humidified atmosphere containing 5% CO₂. The use of human PBMCs was specifically approved by The Medical Ethics Committee of Shanghai Blood Center, (Shanghai, China). Prior to donating blood, the volunteers were informed and provided written informed consent for the scientific research use of blood samples.

Co-culture of PBMCs with or without tumor cells. A total of 2x10^5 MCF-7, MDA-MB-231, A549, A375, HeLa, or HepG2 cells were seeded per a well in their respective growth medium for 30 min and then 2x10^4 PBMCs were added to each well. Tumor cell conditioned media was collected from the cultures after 24 h. PBMCs were exposed to tumor cell conditioned media. After 30 min, anti-CD3 (1 µg/ml) and anti-CD28 (1 µg/ml) were added to the tumor cells/PBMCs co-culture system. After 48 h of co-culture, cell culture supernatants were collected and analyzed for IL-2 and IFN-γ using the HTRF kit (Cisbio; PerkinElmer, Inc.) according to the manufacturer's protocol.

Reverse transcription-quantitative PCR. Total RNA was extracted using TRIzol® reagent according to the manufacturer's protocol (Invitrogen; Thermo Fisher Scientific, Inc.). RNA (1 µg) was used to synthesize cDNA using a PrimeScript RT Reagent kit (Takara Bio, Inc.) according to the manufacturer's protocol. qPCR was performed for PD-L1, PD-L2, CD80, CD86, herpesvirus entry mediator (HVEM), CD70, CD137, OX40L and GAPDH. The sequences of the primer pairs used are shown in Table I. The thermocycling conditions were: 95°C for 10 min; followed by 40 cycles of 95°C for 30 sec, 60°C for 30 sec and 72°C for 30 sec. The fold changes of each gene were calculated using the ΔΔCq (quantification cycle) method, and gene expression levels were normalized to GAPDH (29).

Table I. Forward and reverse primers used for all RT-qPCR analyses.

<table>
<thead>
<tr>
<th>Gene</th>
<th>Forward</th>
<th>Reverse</th>
</tr>
</thead>
<tbody>
<tr>
<td>GAPDH</td>
<td>AGCCGCATCTCTTTTGGGTTGA</td>
<td>TGACGAAATGCGGCGGTCA</td>
</tr>
<tr>
<td>PDL1</td>
<td>GCTGCATAATTTGTCTGTGCTG</td>
<td>AATTCGCTTAGTCTGCGGACC</td>
</tr>
<tr>
<td>IDO1</td>
<td>GGCTGTTGGAATGGCTTCTC</td>
<td>ATGTGCGCTGCTTCTGGG</td>
</tr>
<tr>
<td>PDL2</td>
<td>CAGCAATGTCCTGAGCTGGAAT</td>
<td>GGAACGTGATGCTCAGGACC</td>
</tr>
<tr>
<td>TIM3</td>
<td>GAATTACAGAGGGAAGTCC</td>
<td>AGGGACATCTCCCTGCTGGG</td>
</tr>
<tr>
<td>LAG3</td>
<td>ACCCATCCTCAGGAGGTAA</td>
<td>GTCGCCACTGTCTTCTCTCAA</td>
</tr>
<tr>
<td>CTLA4</td>
<td>CCGTGCACAGAATGCGTCT</td>
<td>ACATCGTCTGCTTCTCTGTTC</td>
</tr>
<tr>
<td>CD80</td>
<td>TCTGTTCAGGCTGTCTCGTGTA</td>
<td>TGGTTCTGGTCTGACATCC</td>
</tr>
<tr>
<td>CD86</td>
<td>ATTCGACAGTCTGAGCGGAGT</td>
<td>CCAAGGAAATGCTGCTGGG</td>
</tr>
<tr>
<td>CD28</td>
<td>ACACCTTTCTGACAAGGTCCC</td>
<td>AGCAGTGCTGCTTCTTACTCC</td>
</tr>
<tr>
<td>ICOS</td>
<td>TTAGACACTGAAACGCGGAGA</td>
<td>AAAACTGCGCAACGTTCTTC</td>
</tr>
<tr>
<td>HVEM</td>
<td>GCTTTGAGCGTGGCTGCTGTA</td>
<td>TGGTCTGCTGCTGACATCC</td>
</tr>
<tr>
<td>CTLA</td>
<td>GACCCCTCAAGGACAGGTGG</td>
<td>TTCTCGGACGACAGAAGG</td>
</tr>
<tr>
<td>CD137L</td>
<td>CGCAGTCTCTCGTCATGGA</td>
<td>CCTCTTTTGTAGCTGACGCGCC</td>
</tr>
<tr>
<td>CD70</td>
<td>GACACACTCTCGACACAACCT</td>
<td>TAATGACAGACAGGTGCGG</td>
</tr>
<tr>
<td>OX40L</td>
<td>AGGCCAAAGATTGCAGAGGGA</td>
<td>CAGTGCTGACATCTTACCTGAA</td>
</tr>
</tbody>
</table>
FACS of tumor cells. Cells were incubated with allophycocyanin-conjugated anti-human-CD274 antibody (1:100) at 4°C for 30 min in the dark for flow cytometry analysis using a Guava® easyCyte Benchtop flow cytometer and FlowJo software (FlowJo™; version 10.6.1; FlowJo LLC) was used to analyze the data.

Immunoblot analysis. MCF-7 and MDA-MB-231 cells were plated in 6 well plates at a density of 1x10^5 cells/ml. The cells were co-cultured with Jurkat cells (2x10^5) in serum-free medium, and treated with anti-CD3 (1 µg/ml) and anti-CD28 (1 µg/ml) for 5, 15 and 30 min, respectively. After the treatment, Jurkat cells were washed in PBS and lysed with RIPA lysis buffer (CoWin Biosciences). Protein concentrations were determined using a bicinchoninic acid assay kit (Thermo Fisher Scientific, Inc.). Western blot analysis was performed as previously described (30). Rabbit polyclonal antibodies against human phospho-AKT, AKT, phospho-ERK, ERK and GAPDH were used at a dilution of 1:1,000 at 4°C in the dark overnight. A horse radish peroxidase-conjugated secondary goat antibody against rabbit immunoglobulin G was used at a dilution of 1:5,000 at room temperature for 1 h. Signals were visualized using Pierce Western Blotting Substrate Plus (Thermo Fisher Scientific, Inc.) and a ChemiDoc XR® system (Bio-Rad Laboratories, Inc.).

Statistical analysis. Data were analyzed with GraphPad Prism version 5.0 (GraphPad Software, Inc.). The results were analyzed using a two-way ANOVA followed by post-hoc Newman-Keuls test. All data are presented as the mean ± standard error of the mean. P<0.05 was considered to indicate a statistically significant difference.

Results

Tumor cells inhibit the activation of Jurkat cells. NFATs are a family of transcription factors which serve important roles in the immune response (31). The human T lymphocyte-based Jurkat cell line expressing luciferase gene under the control of NFAT response elements can be used to study NFAT activation following various stimuli (32). To investigate the effects of the MCF-7, MDA-MB-231, A549, A375, HeLa and HepG2 cells on immune cell activation, these cells were co-cultured with Jurkat cells which were transfected with PGL3-NFAT-TA-Luciferase plasmid and stimulated with anti-CD3 and anti-CD28 antibodies. The results showed that MCF-7, MDA-MB-231 or HeLa cells significantly inhibited the anti-CD3/CD28-induced expression of luciferase in Jurkat cells (Fig. 1A), whereas the other tumor cells did not result in significant changes. However, the conditioned media collected from any of the cell lines, including MCF-7, MDA-MB-231 or HeLa cells, did not significantly affect luciferase expression in Jurkat cells (Fig. 1B). These results suggest that the direct interaction between tumor cells and Jurkat cells, rather than the factors secreted by the tumor cells, inhibited Jurkat cell activation.

Tumor cells inhibit cytokine secretion from PBMCs. Jurkat is an immortalized cell line of human T lymphocytes (13).
Figure 2. Expression of immune checkpoint markers in various cancer cell lines. mRNA expression levels of (A) PD-L1, (B) PD-L2, (C) CD80, (D) CD86, (E) HVEM, (F) CD70, (G) CD137 and (H) OX40L in tumor cell lines. *P<0.05, **P<0.01, ***P<0.001. (I) mRNA expression levels of immune checkpoint receptors in Jurkat cells. Gene expression was normalized to GAPDH in the same sample. (J) FACS analysis of PD-L1 in various cancer cell lines. APC-conjugated anti-human-PD-L1 was used as a binding antibody to cell-surface PD-L1 protein (red line, isotype control staining; blue line, PD-L1 staining). Cell count has been normalized to the peak height at the mode of the distribution, such that absolute count is presented as a percent of the total count. (K) Quantification of the FACS analysis shown in (J). *P<0.01, **P<0.001 vs. 293. APC, allophycocyanin.
Primary PBMCs isolated from whole blood samples were used to investigate the effects of tumor cells. PBMCs were co-cultured with various tumor cell lines and stimulated with anti-CD3/CD28. The secretion of IFN-γ and IL-2 was measured. The results showed that MCF-7, MDA-MB-231 and A549 cells significantly inhibited IFN-γ and IL-2 secretion from PBMCs (Fig. 1C). However, the conditioned media collected from any of the cell lines, including MCF-7, MDA-MB-231 or A549 cell cultures had no effect on cytokine secretion from PBMCs (Fig. 1D). Therefore, similar to the Jurkat cells, a direct interaction between tumor cells and PBMCs resulted in the suppression of cytokine secretion.

Expression of immune checkpoint markers in various tumor cell lines. The aforementioned results suggest that different tumor cells have different effects on suppressing immune cell function. Thus, whether the expression levels of immune checkpoint proteins on tumor cells affected their immune-suppressive activity was determined. The mRNA expression levels of PD-L1 and other immune checkpoint genes, PD-L2, CD80, CD86, HVEM, CD70, CD137 and OX40L were measured in these tumor cells (Fig. 2A-H). RT-qPCR analysis demonstrated that Hela, MCF-7 and MDA-MB-231 cells expressed significantly high levels of PD-L1 compared with 293 cells (P<0.01; Fig. 2A). PD-L1 and PD-L2 expression was significantly higher in A375 and HeLa cells (P<0.01; Fig. 2A and B, respectively); and the expression of HVEM was significantly high in MDA-MB-231 and HeLa cells (P<0.05; Fig. 2A and B). Notably, PD-1 was highly expressed in Jurkat cells compared with the other immune checkpoint receptors, although this was not significant (Fig. 2I). FACS analysis also confirmed that the protein expression levels of PD-L1 were considerably higher in MCF-7 and MDA-MB-231 compared to other tumor cells (Fig. 2J and K), consistent with the higher mRNA expression levels in these cells. Taken together, these findings suggest that the immune-suppressive activity of tumor cells may be associated with the expression levels of PD-L1 in these cells.

**PD-1/PD-L1 inhibitor restores the function of lymphocytes.** MCF-7 and MDA-MB-231 cells had the highest level of PD-L1 expression, and the largest inhibitory effect on T-cell activation and cytokine secretion in the co-culture system. Therefore, the two tumor cell lines were used to investigate the effect of the PD-1/PD-L1 inhibitor on the function of lymphocytes. Stimulation of Jurkat cells with anti-CD3 and anti-CD28 antibodies significantly induced the expression of luciferase, whereas MCF-7 and MDA-MB-231 cells significantly inhibited the expression of luciferase (Fig. 3A). In the present study, A0-L, a small molecular inhibitor of the PD-1/PD-L1 interaction was used. A0-L significantly restored the expression of luciferase in the tumor-Jurkat cell co-culture (Fig. 3A).

Similar results were observed in the PBMC cytokine secretion assay. Anti-CD3 and anti-CD28 co-stimulation induced IFN-γ and IL-2 secretion from PBMCs, which was significantly inhibited by the co-culture with MCF-7 and MDA-MB-231 cells (Fig. 3B and C). Blocking the PD-1/PD-L1 interaction between PBMCs and tumor cells with A0-L significantly increased the secretion of IFN-γ and IL-2 (Fig. 3B and C). Therefore, these results showed that blocking the PD-1/PD-L1 interaction effectively abrogated the inhibition of immune cell functions by tumor cells.

**Tumor cells with high PD-L1 expression suppress pathways involved in T cell activation.** MCF-7 or MDA-MB-231 cells were co-cultured with Jurkat cells to investigate the regulatory effects of tumor cells on AKT and ERK1/2 phosphorylation in immune cells. Jurkat cells were stimulated with anti-CD3 and anti-CD28 (1 µg/ml) antibodies for various durations, and AKT and ERK1/2 phosphorylation...
was assessed by western blotting. Anti-CD3 and anti-CD28 were increased phosphorylation of AKT and ERK1/2 in a time-dependent manner (Fig. 4A and B). By contrast, Jurkat cells co-cultured with MCF-7 or MDA-MB-231 cells had significantly reduced phosphorylation levels of AKT and ERK1/2 following anti-CD3 and anti-CD28 stimulation (Fig. 4A and B). These results indicate that the tumor cells with high PD-L1 expression suppressed the pathways involved in T-cell activation, such as the CD28-PI3K-AKT and TCR-ZAP70-RAS-ERK pathways. Additionally, treatment with A0-L suppressed the inhibitory effects of MCF-7 cells on the CD28-PI3K-AKT and TCR-ZAP70-RAS-ERK pathways in Jurkat cells and restored phosphorylation levels of AKT and ERK1/2 (Fig. 4C and D).

Discussion

Previous findings have shown that PD-L1 is expressed on the surface of tumor cells in a number of different types of cancer and could induce immunosuppression to enable the host to evade antitumor immune responses (33,34). PD-L1, as an immunosuppressive factor and the receptor of PD-L1, is a critical negative regulator of cancer biology with the capacity to support cancer development, growth, invasion and metastasis (35). PD-L1 has
also been established studied as a biomarker of a number of different types of cancer, and several studies demonstrated that PD-L1 expression may be used to predict the outcome of the disease. In the present study, the results showed that PD-L1 mRNA expression levels were upregulated in MCF-7 and MDA-MB-231 tumor cells, consistent with the high protein expression levels of PD-L1 in these cells. Additionally, MCF-7 and MDA-MB-231 cells significantly downregulated T-cell activity and cytokine secretion, which was associated with the upregulated expression of PD-L1 in these tumor cells. Blocking PD-1/PD-L1 interaction with A0-L, a PD-1/PD-L1 inhibitor, significantly restored the activation of Jurkat cells and the secretion of IFN-γ and IL-2 from PBMC cells, which were significantly inhibited by MCF-7 and MDA-MB-231 cells. The results suggest that PD-L1 is upregulated in specific tumor cells and may downregulate T-cell activity by binding to the PD-1 receptor on T cells.

T-cell activation is initiated by the binding of TCRs to their physiological ligands, which are foreign peptides bound to the MHC expressed on antigen-presenting cells (APCs) (36). Upon activation of the TCR, the Src family kinase tyrosine-protein kinase Lck (LCK) becomes activated. The activated LCK phosphorylates CD3 chains, which promote the recruitment and subsequent activation of another tyrosine kinase, ZAP-70, and recruitment of a number of other protein kinases involved in the activation of different signaling cascades, such as RAS and ERK (37,38). CD28 is a co-stimulatory molecule that promotes T-cell activation (39). Upon ligand binding, CD28 recruits and activates PI3K, which in turn activates AKT by phosphorylation (40). The CD28-PI3K-AKT and TCR-ZAP70-RAS-ERK pathways are two major functional signaling pathways involved in T cell activation. When the TCR engages with an antigen peptide and MHC, the T cells are activated via signal transduction, and the primary signaling pathway involved is the TCR-ZAP70-RAS-ERK pathway (41,42). Co-ligation of other cell surface receptors provides additional signals required to enhance T cell activation. CD28 is a costimulatory molecule that promotes T cell proliferation, cytokine production, cell survival and cellular metabolism. CD28-PI3K-AKT is the primary downstream signaling pathway of CD28 (37,38).

In the present study, it was demonstrated that tumor-T cell interaction through PD-1/PD-L1 significantly inhibited the above-mentioned pathways. Tumor cells with high PD-L1 expression inhibited TCR-dependent ERK phosphorylation and CD28-dependent AKT phosphorylation. These results suggest that PD-L1 mediates its inhibitory effects on T cell activation by regulating TCR signaling and CD28 signaling.

The PD-1/PD-L1 targeting strategy was a breakthrough in immunotherapy that restores the functions of T cells (e.g., immune cell activation and differentiation, and cytokine secretion) and promotes immune response (43-46). Keytruda and Tecentriq have been approved by the FDA, their mechanisms of action are well understood, and their clinical efficacy and pharmacodynamics data have been determined. Although several PD-1/PD-L1 inhibitors have been approved for cancer therapy, they are more effective in treating certain tumors over others (47). The results of the present study suggest that tumor cells with a higher expression level of PD-L1 may have higher immunosuppressive activity, and drugs targeting the PD-1/PD-L1 interaction may have improved therapeutic effects on tumors with higher expression levels of PD-L1.

Acknowledgements

Not applicable.

Funding

The present study was supported by grants from The National Science & Technology Major Project: Key New Drug Creation and Manufacturing Program (Beijing, China; grant no. 2017ZX09101004-012-008); Personalized Medicines-Molecular Signature-based Drug Discovery and Development; Strategic Priority Research Program of the Chinese Academy of Sciences (Beijing, China; grant no. XDA12040212), and Shanghai Commission of Science and Technology (Shanghai, China; grant no. 16431901500).

Availability of data and materials

The datasets used and/or analyzed during the present study are available from the corresponding author on reasonable request.

Authors’ contribution

YZ performed the majority of the experiments, analyzed the data and wrote the manuscript. YCF performed some of the experiments. JL supervised the study, analyzed the data and wrote the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The use of human PBMCs was specifically approved by The Medical Ethics Committee of Shanghai Blood Center, (Shanghai, China). Prior to donating blood, the volunteers were informed and provided written informed consent for the scientific research use of blood samples. Due to ethical constraints, no clinical information on the blood donors was obtained.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

References


12. Janssen LME, Ramsay EE, Logsdon CD and Overwijk WW: JAK3/STAT5 inhibition reduces tumor cell proliferation but not on tumor cells, is a favorable prognostic factor for head 


