

Communication

Establishment of Anti-Dog PD-L1 Monoclonal Antibodies for Immunohistochemistry

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Abstract: Immune checkpoint blockade therapy has shown successful clinical outcomes in multiple human cancers. In dogs, several types of tumors resemble human tumors in many respects. Therefore, several groups have developed the anti-dog programmed cell death ligand 1 (dPD-L1) monoclonal antibodies (mAbs) and showed efficacy in several canine tumors. To examine the abundance of dPD-L1 in canine tumors, anti-dPD-L1 diagnostic mAbs for immunohistochemistry are required. In this study, we immunized the peptide in the dPD-L1 intracellular domain, and established anti-dPD-L1 mAbs, L₁Mab-352 (mouse IgG₁, kappa) and L₁Mab-354 (mouse IgG₁, kappa). In enzyme-linked immunosorbent assay, L₁Mab-352 and L₁Mab-354 showed high binding affinity to the dPD-L1 peptide, and the dissociation constants (K_D) were determined as 6.9×10^{-10} M and 7.2×10^{-10} M, respectively. Furthermore, L₁Mab-352 and L₁Mab-354 were applicable for the detection of dPD-L1 in immunohistochemical analysis in paraffin-embedded dPD-L1-expressed cells. These results indicated that L₁Mab-352 and L₁Mab-354 are useful for detecting dPD-L1 in immunohistochemical analysis.

Keywords: dPD-L1; monoclonal antibody; peptide immunization; immunohistochemistry

1. Introduction

Immune checkpoint blockade therapy has recently revolutionized the treatment and clinical outcome of several cancer types [1]. The therapies targeting programmed cell death 1 (PD-1), programmed cell death ligand 1 (PD-L1), and cytotoxic T-lymphocyte-associated antigen 4 (CTLA-4) have become the standard therapy for several human tumors. The therapeutic mAbs against PD-1 receptor (nivolumab and pembrolizumab), PD-L1 (atezolizumab, durvalumab, and avelumab), CTLA-4 (tremelimumab and ipilimumab) were approved by Food and Drug Administration (FDA) [2]. These mAbs have been used in either monotherapy or combinatorial therapy with chemotherapy, radiation therapy, or other modalities [1]. These antibodies improve the response rates, progression-free survival, and overall survival in patients with various types of cancers [3–5].

PD-L1 (CD274) is a type I membrane protein expressed on non-lymphoid cells and a ligand of PD-1 [6,7]. PD-1 plays a critical role in the negative regulation of immune cells including cytotoxic T lymphocyte (CTL) and is important for antitumor immunity. CTLs are activated by recognizing presented antigens by T cell receptor (TCR)-CD3 complex [8]. After the recognition, zeta-chain associated protein kinase (ZAP-70) is recruited to the TCR-CD3 complex and phosphorylated by lymphocyte protein tyrosine kinase, which transduces the downstream signaling [9,10]. In contrast, SH2-containing protein tyrosine phosphatase-2 (SHP-2) is recruited to PD-1 stimulated with PD-L1 [11]. Since SHP-2 dephosphorylates phosphorylated ZAP-70, PD-1 can inhibit the activation signals in CTLs.

With the increase in life span in both humans and dogs, the cancer incidence has increased as well [12]. Several naturally occurring tumors in dogs resemble human tumors in many respects. Therefore, the research on canine tumor therapy can generate knowledge that informs and prioritizes new tumor therapy in humans.

The development of dog PD-L1 (dPD-L1) mAbs has been reported [13–17]. Using a canine chimeric mAb targeting PD-L1, tumor regression of undifferentiated sarcoma and oral melanoma was achieved [14,16,17]. On the other hand, tumors that did not respond to the treatment also existed. Furthermore, the combination therapy of anti-dPD-L1 mAbs with hypofractionated radiotherapy is more effective to prolong overall survival [18]. Therefore, the development of anti-dPD-L1 mAbs for diagnostic use is essential for the improvement of efficacy.

In this study, we established anti-dog PD-L1 (dPD-L1) mAbs (L₁Mab-352 and L₁Mab-354) by peptide immunization and showed the usefulness of the mAbs for immunohistochemical analysis in paraffin-embedded PD-L1-positive cells.

2. Materials and Methods

2.1. Preparation of cell lines

Chinese hamster ovary (CHO)-K1 and P3X63Ag8U.1 (P3U1) cells were obtained from the American Type Culture Collection (Manassas, VA).

The synthesized DNA (Eurofins Genomics KK, Tokyo, Japan) encoding signal sequence of N-terminus (1-MRMFSVFTFMAYCHLLKA-18) deleted dPD-L1 (dPD-L1, Accession No.: NM_001291972) was subsequently subcloned into a pCAGzeo_ssPA16 vector (IL2-signal sequence and PA16 tag added to N-terminus of construct). The amino acid sequence of the tag system was as follows: PA16 tag [19–22], sixteen amino acids (GLEGGVAMPGAEDDVV). The PA16 tag can be detected by an anti-human podoplanin mAb (clone NZ-1) [19–35]. The dPD-L1 plasmid was transfected into CHO-K1 cells, using a Neon transfection system (Thermo Fisher Scientific Inc., Waltham, MA). Stable transfectants were established through cell sorting using a cell sorter (SH800; Sony Corp., Tokyo, Japan), after which cultivation in a medium, containing 0.5 mg/mL of Zeocin (InvivoGen, San Diego, CA) was conducted.

CHO-K1, PA16-dPD-L1-overexpressed CHO-K1 (CHO/dPD-L1) and P3U1 cells were cultured in a Roswell Park Memorial Institute (RPMI)-1640 medium (Nacalai Tesque, Inc., Kyoto, Japan), with 10% heat-inactivated fetal bovine serum (FBS; Thermo Fisher Scientific Inc.), 100 units/mL of penicillin, 100 µg/mL of streptomycin, and 0.25 µg/mL of amphotericin B (Nacalai Tesque, Inc.). All cells were grown in a humidified incubator at 37°C, in an atmosphere of 5% CO₂ and 95% air.

2.2. Production of hybridomas

A five-week-old BALB/c mouse was purchased from CLEA Japan (Tokyo, Japan). The animal was housed under specific pathogen-free conditions. All animal experiments were approved by the Animal Care and Use Committee of Tohoku University (Permit number: 2022MdA-001). The dPD-L1 peptide (260-KKHGRMMDVEKC-271) and keyhole limpet hemocyanin-conjugated dPD-L1 peptide (dPD-L1 peptide-KLH) were purchased from Eurofins Japan.

To develop mAbs against dPD-L1, we intraperitoneally immunized one mouse with the dPD-L1 peptide-KLH (100 µg) plus Imject Alum (Thermo Fisher Scientific, Inc.). The procedure included three additional injections every week (100 µg), which were followed by a final booster intraperitoneal injection (100 µg), two days before harvesting splenocytes. The harvested splenocytes were subsequently fused with P3U1 cells, using PEG1500 (Roche Diagnostics, Indianapolis, IN). For the hybridoma selection, cells were cultured in the RPMI-1640 medium with 10% FBS, 100 units/mL of penicillin, 100 µg/mL of streptomycin, 0.25 µg/mL of amphotericin B, 5 µg/mL of Plasmocin, 5% Briclone (NICB, Dublin, Ireland), and hypoxanthine, aminopterin and thymidine (HAT; Thermo Fisher Scientific, Inc.). The supernatants were subsequently screened using enzyme-linked immunosorbent assay (ELISA) using the dPD-L1 peptide.

2.3. Purification of mAbs

The cultured supernatants of L₁Mab-352 and L₁Mab-354-producing hybridomas were filtrated with Steritop (0.22 µm, Merck KGaA, Darmstadt, Germany). The filtered supernatants were subsequently applied to 1 mL of *Ab-Capcher ExTra* (ProteNova, Inc., Kagawa, Japan). After washing with phosphate-buffer saline (PBS), bound antibodies were eluted with an IgG elution buffer (Thermo Fisher Scientific, Inc.), followed by immediate neutralization of eluates, using 1M Tris-HCl (pH 8.0). Finally, the eluates were concentrated, after which PBS was replaced with the elution buffer using Amicon Ultra (Merck KGaA).

2.4. ELISA

The dPD-L1 peptide was immobilized on Nunc Maxisorp 96-well immunoplates (Thermo Fisher Scientific, Inc.) at 1 µg/mL for 30 minutes at 37°C. After washing with PBS containing 0.05% Tween20 (PBST; Nacalai Tesque, Inc.), wells were blocked with 1% bovine serum albumin (BSA) in PBST for 30 minutes at 37°C. Then, plates were incubated at 1 µg/mL of L₁Mab-352 and L₁Mab-354, followed by peroxidase-conjugated anti-mouse immunoglobulins (1:1000; Agilent Technologies, Inc., Santa Clara, CA). Finally, enzymatic reactions were conducted using the ELISA POD substrate TMB kit (Nacalai Tesque, Inc.) and stopped by the addition of 1M H₂SO₄. The absorbance at 450 nm was measured by using an iMark microplate reader (Bio-Rad Laboratories, Inc., Berkeley, CA).

To determine the dissociation constant (K_D), L₁Mab-352 and L₁Mab-354 were serially diluted from 40 µg/mL to 2.4 ng/mL. The K_D was calculated by fitting saturation binding curves to the built-in; one-site binding models in GraphPad PRISM 8 (GraphPad Software, Inc., La Jolla, CA).

2.5. Flow cytometric analysis

CHO-K1 and CHO/dPD-L1 cells were harvested after a brief exposure to 0.25% trypsin and 1 mM ethylenediaminetetraacetic acid (EDTA, Nacalai Tesque, Inc.). The cells were subsequently washed with 0.1% BSA in PBS and treated with 1 µg/mL NZ-1 for 30 min at 4°C. The cells were treated with 2 µg/mL Alexa Fluor 488-conjugated anti-rat IgG (Cell Signaling Technology, Inc., Danvers, MA). The fluorescence data were collected using the EC800 Cell Analyzer (Sony Corp.).

2.6. Immunohistochemical analysis of paraffin-embedded CHO/dPD-L1

Cell blocks were produced using iPCell (Genostaff Co., Ltd., Tokyo, Japan). To deparaffinize, rehydrate and retrieve antigen, the sections were autoclaved in EnVision FLEX Target Retrieval Solution, High pH (high pH; #DM828, Agilent Technologies, Inc.) at 121°C for 20 minutes. Then, sections were blocked using the Super Block T20 (PBS) Blocking Buffer (Thermo Fisher Scientific, Inc.), incubated with 50 µg/mL L₁Mab-352 or L₁Mab-354 for 1 h at room temperature, and treated with the Envision + Kit for mouse (Agilent Technologies, Inc.) for 30 min. Finally, color was developed using 3,3'-diaminobenzidine tetrahydrochloride (DAB; Agilent Technologies, Inc.) for 5 min, and counterstaining was performed using hematoxylin (FUJIFILM Wako Pure Chemical Corporation, Osaka, Japan).

To inhibit L₁Mab-352 and L₁Mab-354 binding, dPD-L1 peptide (final concentration: 20 µg/mL) was mixed with L₁Mab-352 or L₁Mab-354 (final concentration: 50 µg/mL). After incubation for 1.5 h at room temperature, immunohistochemical analysis was performed.

3. Results

3.1. Establishment of novel dPD-L1 mAbs.

For this study, we immunized a mouse with a KLH-conjugated synthetic peptide corresponding to an intracellular region of dPD-L1. To produce hybridomas, the splenocytes from the mouse were fused with P3U1 cells by using polyethylene glycol. The wells which reacted with the synthetic peptide were selected by ELISA. After limiting dilution and additional screening, two clones L₁Mab-352 (mouse IgG₁, kappa) and L₁Mab-354 (IgG₁, kappa) were finally established (Figure 1).

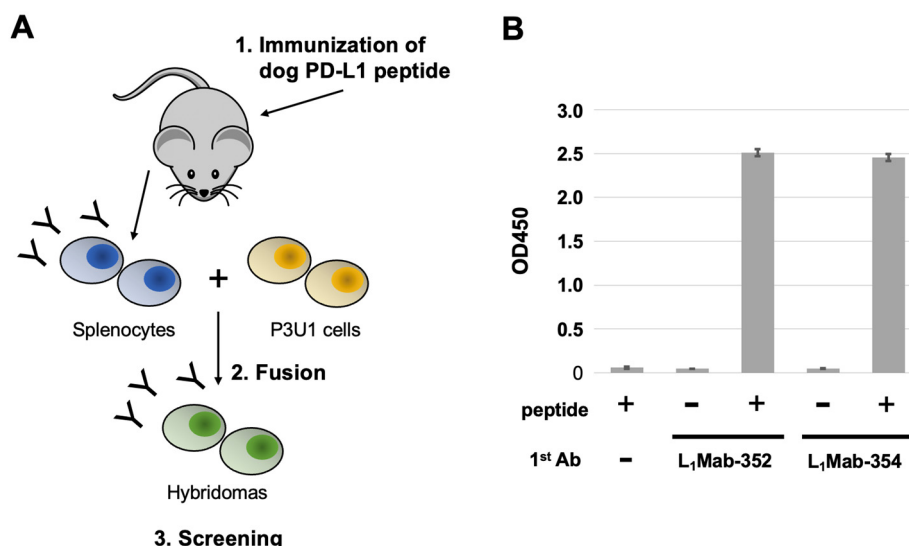


Figure 1. Establishment of novel dPD-L1 mAbs, L₁Mab-352 and L₁Mab-354. (A) The workflow of the establishment of L₁Mab-352 and L₁Mab-354. First, dPD-L1 peptide conjugated with KLH was immunized into a mouse. The splenocytes were then fused with P3U1 cells. After six days, hybridoma supernatants were selected by ELISA. (B) Synthesized peptides of dPD-L1 peptide (1 µg/mL) were immobilized on immunoplates for 30 min at 37°C. The plates were incubated with 1 µg/mL of L₁Mab-352 and L₁Mab-354, followed by the treatment of peroxidase-conjugated anti-mouse immunoglobulins. Optical density was measured at 450 nm (OD₄₅₀) using a microplate reader. Error bars represent means ± SDs (n = 3).

3.2. Kinetic analyses of L₁Mab-352 and L₁Mab-354 against the dPD-L1 peptide.

To determine the K_D of L₁Mab-352 and L₁Mab-354 with the dPD-L1 peptide, we conducted the kinetic analysis using ELISA. After immobilization of the dPD-L1 peptide on immunoplates, serially diluted L₁Mab-352 and L₁Mab-354 were added. The mean of the absorbance at 450 nm was plotted versus the concentrations of L₁Mab-352 and L₁Mab-354. The K_D values of L₁Mab-352 and L₁Mab-354 for the dPD-L1 peptide were calculated as 6.9×10^{-10} M and 7.2×10^{-10} M, respectively (Figure 2).

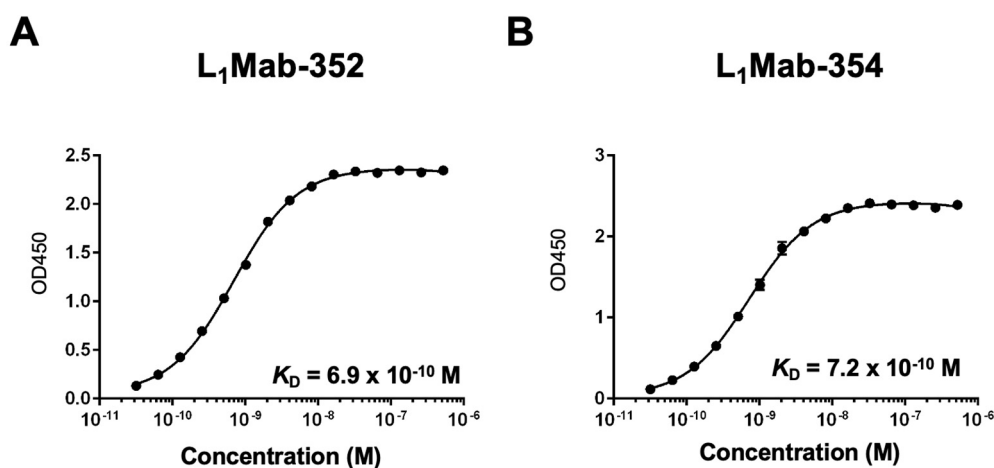


Figure 2. The determination of the binding affinity of L₁Mab-352 and L₁Mab-354 by ELISA. Synthesized peptides of dPD-L1 peptide (1 µg/mL) were immobilized on immunoplates for 30 min at 37°C. The plates were incubated with L₁Mab-352 (A) and L₁Mab-354 (B) at 2.4 ng/mL–40 µg/mL, followed by the treatment of peroxidase-conjugated anti-mouse immunoglobulins. Optical density was measured at 450 nm (OD₄₅₀) using a microplate reader. The dissociation constant (K_D) was calculated using GraphPad PRISM 8. Error bars represent means ± SDs (n = 3).

3.3. Immunohistochemical analysis of L₁Mab-352 and L₁Mab-354 using paraffin-embedded CHO/dPD-L1.

We next performed immunohistochemical analysis against paraffin-embedded CHO/dPD-L1 cells (Figure 3A) using L₁Mab-352 and L₁Mab-354. We first confirmed the cell surface expression of dPD-L1 using flow cytometry (Figure 3B). Then, we prepared the paraffin-embedded CHO-K1 and CHO/dPD-L1 cells, and stained the section using L₁Mab-352 and L₁Mab-354. As shown in Figure 3C,D, L₁Mab-352 and L₁Mab-354 stained to the section of CHO/dPD-L1, but not that of CHO-K1. These results indicated that L₁Mab-352 and L₁Mab-354 can stain dPD-L1 in paraffin-embedded samples.

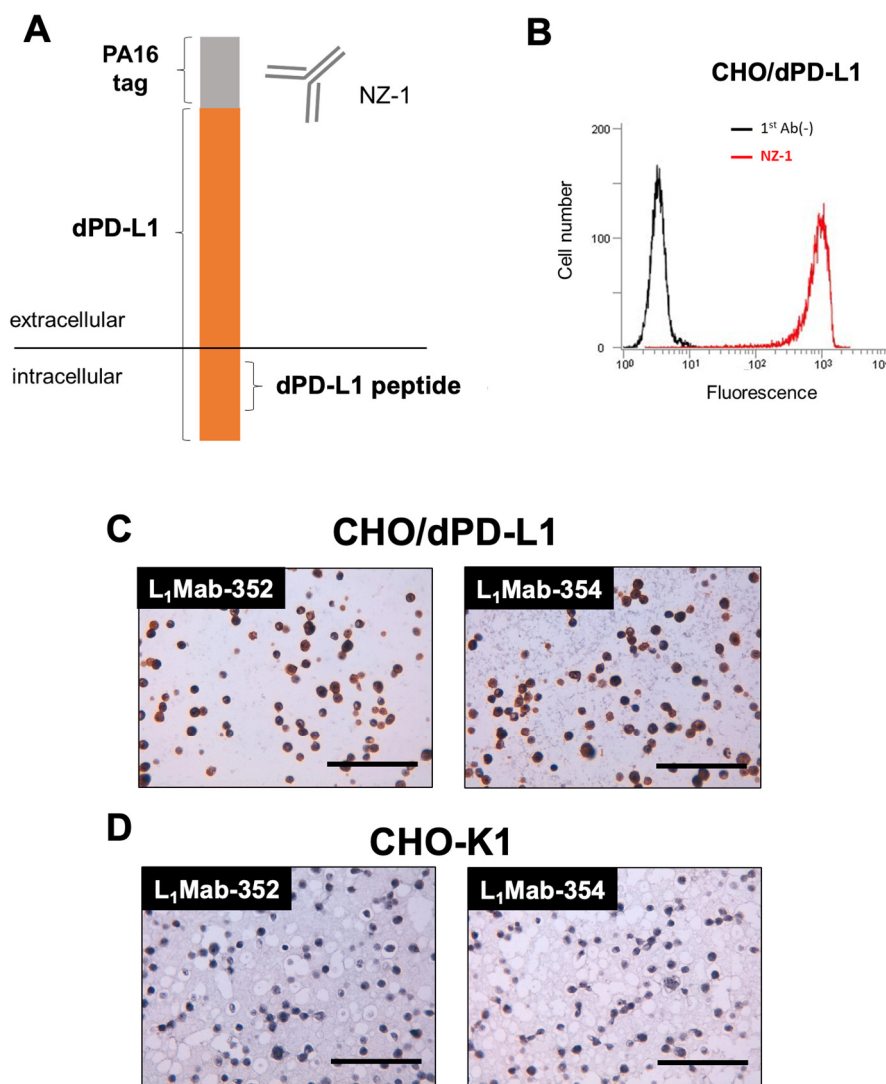


Figure 3. Immunohistochemical staining of paraffin-embedded CHO/dPD-L1 with L₁Mab-352 and L₁Mab-354. (A) The construction of dPD-L1. PA16 tag, the epitope of NZ-1, was tagged to the N-terminus of dPD-L1. The dPD-L1 peptide is located in the intracellular domain. (B) The cell surface expression of dPD-L1 was confirmed by flow cytometry using NZ-1. Red line: NZ-1. Black line: negative control. (C,D) Immunohistochemical analysis of paraffin-embedded CHO/dPD-L1 (C) and CHO-K1 (D) using 50 μg/mL of L₁Mab-352 and L₁Mab-354. Scale bar = 100 μm.

3.4. Peptide blocking of L₁Mab-352 and L₁Mab-354 in Immunohistochemical analysis.

To confirm the specificity, we performed peptide blocking in the immunohistochemical analysis. As shown in Figure 4, the reactivity of L₁Mab-352 and L₁Mab-354 was blocked in the presence of dPD-L1 peptide. This result indicated that the signals of staining with L₁Mab-352 and L₁Mab-354 are caused by the specific binding to dPD-L1.

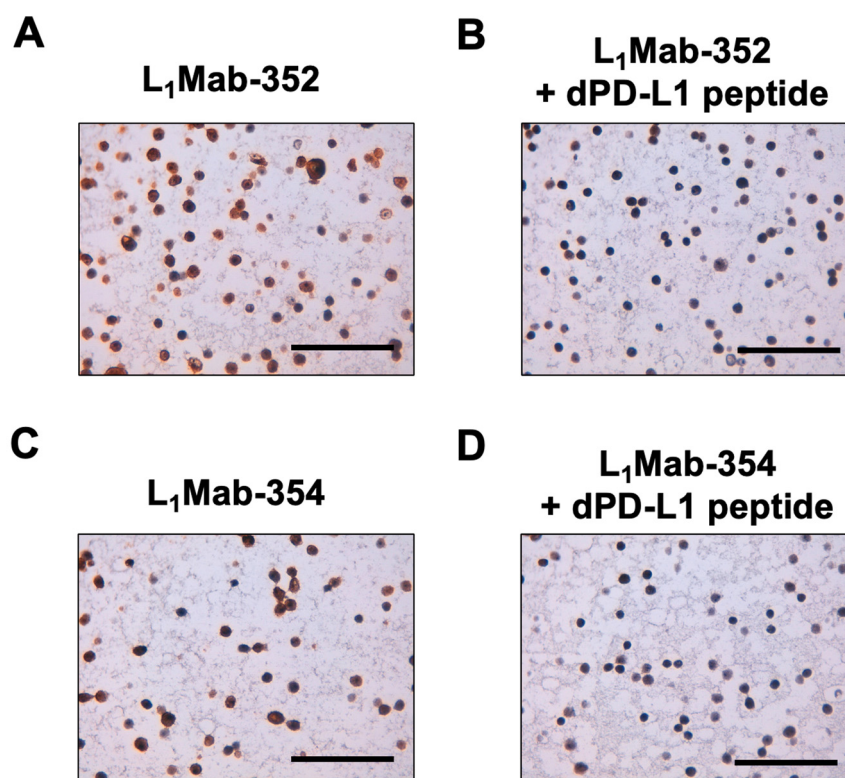


Figure 4. The blockade of L₁Mab-352 and L₁Mab-354 reactivity by dPD-L1 peptide. Immunohistochemical staining of paraffin-embedded CHO/dPD-L1 with L₁Mab-352 (50 µg/mL) (A), L₁Mab-352 (50 µg/mL) plus dPD-L1 peptide (20 µg/mL) (B), L₁Mab-354 (50 µg/mL) (C), and L₁Mab-354 plus dPD-L1 peptide (20 µg/mL) (D). Scale bar = 100 µm.

4. Discussion

Immune checkpoint blockade therapies have improved the patients' outcomes with various types of tumors [36]. However, only 30% of patients receive the benefit from the therapy [37]. Immunohistochemistry analyses to determine the PD-L1-positive tumor cells are widely validated and used as predictive biomarkers to select the patients for immune checkpoint blockade therapy [38]. However, different diagnostic anti-PD-L1 mAb clones were approved for specific therapeutic ones by FDA. For instance, clone 22C3 has been utilized as a predictive biomarker for pembrolizumab in several cancers. A clone 28-8 has been approved as a complementary assay for nivolumab. Each clone has a different cut-off point and cancer-specific scoring algorithm [2]. Both 22C3 and 28-8 recognize the extracellular domain of PD-L1, and exhibit similar subcellular patterns of PD-L1 expression in tumor and immune cells [2]. However, different staining patterns by 22C3 and 28-8 are also reported [39].

In this study, we developed novel dPD-L1 mAbs (L₁Mab-352 and L₁Mab-354) using peptide immunization of the intracellular domain, and showed the usefulness for immunohistochemical analysis in paraffin-embedded dPD-L1-positive cells (Figures 3 and 4). Further studies are essential to show whether L₁Mab-352 and L₁Mab-354 apply to formalin-fixed paraffin-embedded canine tumors. An anti-dPD-L1 mAb (6C11-3A11) which recognizes the extracellular domain of dPD-L1 was reported to apply to immunohistochemistry [17]. The comparison of the staining pattern of mAbs targeting dPD-L1 extracellular and intracellular domains could provide supportive information in human cancer diagnosis.

Currently, several anti-dPD-L1 mAbs have been developed for canine tumor therapy [16,17,40]. Like human tumors, standardization of PD-L1 immunohistochemistry will be required in the future.

Author Contributions: T.O. and T.T. performed the experiments. M.K.K. and Y.K. designed the experiments. T.O., H.S., and Y.K. analyzed the data. T.O. and H.S. wrote the manuscript. All authors read and approved the final manuscript and agreed to be accountable for all aspects of the research in ensuring that the accuracy or integrity of any part of the work is appropriately investigated and resolved. All authors have read and agreed to the published version of the manuscript.

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Institutional Review Board Statement: The animal study protocol was approved by the Animal Care and Use Committee of Tohoku University (Permit number: 2022MdA-001) for studies involving animals.

Informed Consent Statement: Not applicable.

Data Availability Statement: All related data and methods are presented in this paper. Additional inquiries should be addressed to the corresponding authors.

Conflicts of Interest: The authors declare no conflict of interest involving this article.

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