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#### Article

# Development of monoclonal antibodies targeting canine PD-L1 and PD-1 and their clinical relevance in Canine Apocrine **Gland Anal Sac Adenocarcinoma**



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Abstract: Canine apocrine gland anal sac adenocarcinoma (AGASACA) is an aggressive canine tu-23 mor originating from the anal sac glands. Surgical resection, with or without adjuvant chemother-24 apy, represents the standard of care for this tumor, but the outcome is generally poor particularly 25 for tumors diagnosed at an advanced stage. For this reason, novel treatment options are warranted, 26 and few recent reports suggested an activation of the immune checkpoint axis in canine AGASACA. 27 In our study we developed canine-specific monoclonal antibodies targeting PD-1 and PD-L1. Forty-28 one AGASACAs with complete clinical and follow-up information were then analyzed by immuno-29 histochemistry for the expression of the two checkpoint molecules (PD-L1 and PD-1) and the pres-30 ence of tumor infiltrating lymphocytes (CD3 and CD20), evaluated within the tumor bulk (intra-31 tumor) and in the surrounding stroma (peritumor). Seventeen AGASACAs (42%) expressed PD-L1 32 in a range between 5% and 95%. Intratumor lymphocytes were predominantly CD3+ T-cells and 33 positively correlated with the number of PD-1+ intratumor lymphocytes ( $\rho$ =0.36; p=0.02). Peritumor 34 lymphocytes were a mixture of CD3+ and CD20+ cells with a variable PD-1 expression (range 0-35 50%). PD-L1 expression negatively affected the survival only in the subgroup of dogs treated with 36 surgery alone (n=14; 576 vs 235 days). The presence of a heterogeneous lymphocytic infiltrate and 37 the expression of PD-1 and PD-L1 molecules support the relevance of the immune-microenviron-38 ment in canine AGASACAs and the potential value of immune checkpoints as promising therapeu-39 tic targets. 40

Keywords: Dog; Apocrine gland anal sac adenocarcinoma; PD-1; PD-L1, Tumor-infiltrating 41 lymphocytes; Immunohistochemistry. 42

#### 1. Introduction

Apocrine gland anal sac adenocarcinoma (AGASACA) is a relatively rare skin 45 neoplasm, representing approximately 17% of perianal tumors of dogs [1]. It is 46

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characterized by an aggressive biological behavior, with a high rate of metastasis affecting 47 the locoregional lymph nodes and, less frequently, distant sites. The median survival time 48ranges from 1 to 2 years, depending on the clinical stage [1–13]. Prognostic factors include 49 regional lymph node metastasis at the time of diagnosis, tumor and regional lymph node 50 size, histological pattern, necrosis and lymphovascular invasion [14–18]. Conversely, the 51 prognostic significance of hypercalcemia remains controversial and Ki67 proliferation 52 index did not demonstrate a prognostic value [18,19]. Surgical excision represents the best 53 treatment option for AGASACAs without distant metastasis [2]. Chemotherapy and 54 radiotherapy are also employed as adjuvant and/or palliative treatments, and toceranib 55 phosphate showed effective results in prolonging survival time [3-13,20]. 56 Immunotherapeutic approaches have not been explored yet. 57

In several canine cancers, the immune checkpoint programmed death-1 (PD-1) and 58 its ligand PD-ligand 1 (PD-L1), were recently suggested as prognostic markers as well as 59 promising therapeutic targets [21–27]. The interaction between PD-1 and PD-L1 is 60 physiologically involved in the immune response regulation, resulting in the activation of 61 inhibitory signals, responsible of a reduced production of antibodies and cytokines by the 62 immune cells [28]. In oncology, the PD-1/PD-L1 axis activation induced by PD-L1-63 expressing tumor cells interacting with PD-1-expressing tumor-infiltrating lymphocytes 64 (TILs), is one of the most studied immune evasion strategies played by cancer, paving the 65 way for the development of immunotherapeutic approaches based on the blockade of 66 these molecules [28]. Recently, the expression of PD-L1 was investigated by 67 immunohistochemistry (IHC) in various canine tumors, among which 95% of the 68 analyzed AGASACAs (19/20) expressed the immune checkpoint molecule, suggesting the 69 activation of the axis [26]. However, these results were limited to a small number of cases 70 and the association with clinico-pathological data was not considered. 71

In this study, we developed canine-specific monoclonal antibodies targeting PD-1 72 and PD-L1. The expression of the two immune checkpoint molecules was evaluated by 73 IHC in 41 surgically resected AGASACAs. The immunophenotypic characterization of 74 TILs was also performed. Furthermore, the correlation of IHC results with clinico-75 pathological features and follow-up data was investigated, in order to elucidate the 76 prognostic role of the immune microenvironment and to provide new therapeutic 77 insights.

#### 2. Materials and Methods

#### Study population

The AGASACA cases diagnosed and treated at the Veterinary Teaching Hospital 81 (VTH) of the University of Turin, in a spanning period of 10 years (2011-2021) were 82 considered for the study. Only dogs that underwent surgical excision of the primary 83 tumor and of regional lymph nodes when found enlarged on CT scan evaluation, were 84 included in the study, regardless of the administration of adjuvant chemotherapy. Dogs 85 diagnosed with stage IV disease before the surgery or having concurrent severe illness 86 that could significantly reduce the survival time were excluded. 87

For each dog, signalment, primary tumor side, tumor size, clinical signs at 88 presentation, presence of hypercalcemia (ionized calcium >1.45 mmol/L), presence of ileo-89 sacral lymphadenopathy, complete blood analysis, pre- and post-contrast whole body 90 computed tomography, type of treatment (surgery with or without adjuvant 91 chemotherapy), and surgical and post-surgical complications were recorded. Follow-up 92 data were obtained from the follow-up visits at the VTH and/or by telephone contact with 93 owners or referring veterinarians. The disease-free interval (DFI) was calculated as the 94 interval between surgery and local recurrence or metastatic spread, and the survival time 95 (ST) as the time from surgery until death for any cause. 96

Histological analysis

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All surgically removed AGASACAs were histologically confirmed and the following 99 morphological features were recorded [19]: predominant tumor pattern (solid, 100 rosettes/tubules and papillary), necrosis (absent or present), inflammatory infiltration 101 (absent or present), status of surgical margins, lympho-vascular invasion (absent or 102 present), cellular pleomorphism (anisokaryosis and anisocytosis), mitotic count and Ki67 103 index evaluated by IHC. 104

#### Generation of anti-PD-1 and anti-PD-L1 monoclonal antibodies

Anti-PD-1 and anti-PD-L1 monoclonal antibodies were generated by Moravian-107 Biotechnology Ltd. (Brno, Czech Republic). BALB/c mice were hyper-immunized with 108 canine recombinant PD-1 and PD-L1 proteins (with a 6xHis, and Fc tag, respectively; Sino 109 Biological Europe GmbH, Eschborn, Germany). For both targets, splenocytes from hyper-110immunized mice were fused with the non-producing mouse myeloma cell line SP2. 111 Supernatants of selected hybrids were screened using dot-blot on nitrocellulose 112 membrane coated with recombinant canine PD1 (Sino Biological Europe GmbH, 113 Eschborn, Germany) or PD-L1 (KingFisher Biotech Inc, Saint Paul, MN, USA) [29]. 114 Reactive supernatants were further validated by ELISA, western blotting and flow 115 cytometry. 116

#### ELISA – PD-L1

Recombinant PD-L1-FC fusion protein (Sino Biological Europe GmbH, Eschborn, 119 Germany) was diluted in carbonate coating buffer (pH 9.6) at 1 µg/mL and a 96-well plate 120 (Costar) was coated with 50 ng of protein overnight at 4°C. Wells were subsequently 121 washed three times with PBST and blocked with blocking buffer (PBS with 2% BSA) for 1 122 hour at room temperature. Blocking buffer was removed and serial dilutions of antibodies 123 (1:50, 1:100, 1:200, 1:400, 1:800, 1:1600 – starting from 1mg/mL) were added and incubated 124 overnight at 4°C. Wells were then emptied, washed three times with PBST and incubated 125 for 1 hour at room temperature with goat anti-mouse HRP conjugated antibody (Dako) 126 diluted in PBS with 2% BSA at 1:1000. After washing the wells, Pierce ECL (Thermo Fisher 127 Scientific, Oxford, UK) was added to the wells and the plate was read using BioTek 128 Synergy plate reader. Data were subsequently analyzed using GraphPad Prism 9.4 129 (GraphPad, San Diego, CA, USA). 130

#### ELISA - PD-1

Recombinant PD-1-FC fusion protein (Sino Biological) was diluted in carbonate 133 coating buffer (in-house; pH 9.6). A 96-well opaque white plate (Costar) was coated 134 overnight at 4°C with 50ng of protein per well. Wells were subsequently washed four 135 times with 400µl of PBST 0,05% (the same for all the following washes) and blocked with 136 blocking buffer (BB; PBST with 3% BSA) for 90min at room temperature (RT). Blocking 137 buffer was removed and serial dilutions of the antibody (purified stock in PBS) were 138 prepared (1000, 100, 10, 1 ng/mL, equal to 100, 10, 1, 0.1 ng/well) and added to plate, 100µl 139 per well. The plate was incubated for 1h in RT. Wells were emptied, washed, and 140 incubated for 1 hour at room temperature with a rabbit anti-mouse polyclonal HRP 141 conjugated antibody (Dako P0260) diluted in BB at 1:1000. After washing the wells, Pierce 142 ECL (Thermo Fisher) was added to the wells and the plate was read immediately using 143 Fluoroskan Ascent plate reader (Thermo Fisher), Luminescence mode and integration 144 time 300ms. Data was analyzed using GraphPad Prism 9.4 (GraphPad, San Diego, CA, 145 USA). 146

#### Western Blotting

Western blotting (WB) for PD-L1 analysis was performed using recombinant PD-L1-149Fc fusion protein (Sino Biological Europe GmbH, Eschborn, Germany) and cell lysates of150K9TCC, K9TCC-SH (both kindly provided by Prof. Deborah Knapp, Purdue University,151USA), and DH82 (ATCC). K9TCC, K9TCC-SH and DH82. All cell lines were selected152

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based on previous publications, which have shown that the cell lines expressed PD-1 153 and/or PD-L1 [27,30,31]. In addition, the K9TCC and K9TCC-SH were considered 154 having different levels of expression of EGFR [32]. EGFR is known to affect PD-L1 stability 155 through modifications in glycosylation levels, allowing to investigate antibody bindings 156 to different patterns of PD-L1 glycosylation [33]. 157

Thirty micrograms of cell lysates were separated on Novex NuPage 4-12% SDS-158 PAGE gel (Thermo Fisher Scientific, Oxford, UK) and transferred to PVDF membrane 159 using iBlot transfer system (Thermo Fisher Scientific, Oxford, UK). Membranes were then 160 blocked with Intercept® (TBS) Blocking Buffer (Li-cor, Cambridge, UK) and incubated 161 with PD1 1.1 or PD-L1 3.1 antibodies at 1:1000 (PD-1 1.1) or 1:200 (PD-L1 3.1) concentration 162 overnight. After the incubation, membranes were washed and incubated with IRDye® 163 800CW Donkey anti-Mouse (Li-cor, Cambridge, UK). The membrane was subsequently 164 washed with PBST and PBS. The membrane was imaged using Odyssey imaging system 165 (Li-cor, Cambridge, UK). 166

#### *Flow cytometry*

Cell lines used for WB, were also utilized in flow cytometry (FC) experiments. Cells 169 were expanded in DMEM/F12 (K9TCC, K9TCC SH) or DMEM (DH82, OSA31) media 170 (both from Gibco, Thermo Fisher Scientific, Oxford UK) supplemented with 10% fetal 171 bovine serum (FBS) (Gibco, Thermo Fisher Scientific, Oxford UK) and Penicillin-172 Streptomycin (Gibco, Thermo Fisher Scientific, Oxford UK). After reaching confluency, 173 cells were detached using 0.25% Trypsin-EDTA (Gibco, Thermo Fisher Scientific, Oxford 174 UK), counted and 5x10<sup>5</sup> cells were used for each antibody. For all cell lines, PD-L1 175 antibodies 1.1, 2.1 and 3.1 were used at concentration of 1:100 diluted in PBS with 2% BSA 176 and incubated for 1 hour. Subsequently, cells were washed with PBS with 2% BSA and 177 incubated with 1:1000 dilution of FITC conjugated anti-mouse polyvalent 178 immunoglobulin secondary antibody (Sigma, Merck KGaA, Darmstadt, Germany), which 179 binds all antibody subclasses. Cells incubated with secondary antibody were used as 180 controls. PD-1 antibody was directly conjugated to APC fluorophore using a 181 commercially available kit (Abcam, Cambridge, UK) and did not require a secondary 182 antibody. Unstained cells were used as controls for PD-1 antibody. 183

For DH82 cell line, as it expresses Fc receptor, cells were incubated with Fc Receptor184Binding Inhibitor (eBioscience, ThermoFisher Scientific, Oxford,UK) for 30 min. Cells185were then washed, resuspended in staining buffer and flow cytometry was performed186using BD LSR Fortessa flow cytometer (Becton Dickinson, Wokingham, UK). Data were187subsequently analyzed using FlowJo (FlowJo LLC, Ashland, OR, USA).188

In addition a K9TCC transiently overexpressing PD-1 was generated via transfection 189 of this cell line with a custom-build pcDNA3.1 plasmid vector encoding for canine PD-1 190 (GeneArt, ThermoFisher Scientific, Oxford,UK). Cells were transfected with the vector 191 using lipofectamine (Gibco, ThermoFisher Scientific, Oxford, UK) and selected using 192 Hygromycin (ThermoFisher Scientific, Oxford, UK). Transfection was confirmed using 193 flow cytometry as described above. 194

#### Immunohistochemistry

Immunohistochemistry (IHC) was performed on 4 µm thick FFPE sections using 197 VECTASTAIN® Universal Quick HRP Kit, Peroxidase, R.T.U. from Vector Laboratories 198 Inc. (Burlingame, CA), following the manufacturer's instructions. Sections were incubated 199 with 0.3% H<sub>2</sub>O<sub>2</sub> for endogenous peroxidase activity blocking, and heat-induced antigen 200 retrieval was then performed in citrate buffer (pH 6.0) (for anti-PD-L1, anti-PD-1, anti-201 CD20 primary antibodies) or in Tris-EDTA buffer (pH 9.0) (for anti-CD3 primary 202 antibody) at 98°C for 20 minutes. After non-specific antibody binding blocking 203 performed with Normal Horse Serum 2.5%, sections were incubated with the following 204 primary antibodies: anti-CD3 (diluted 1:50; Dako, Nowy Sącz, Poland), anti-CD20 205 (diluted 1:1000; Moravian-Biotechnology) and anti-PD-1 (diluted 1:100) for 2h at room 206

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temperature, and anti-PD-L1 (diluted 1:10) overnight at 4°C. Signal was detected using 207 Vectastain Elite ABC kit and ImmPACT DAB from Vector Laboratories Inc. (Burlingame, 208 CA). In each immunohistochemical run positive controls were included (canine lymph 209 node for anti-CD3 and anti-CD20; canine tonsil for anti-PD-1; canine diffuse large B-cell 210 lymphoma known to have high PD-L1 mRNA expression by RNAscope for anti-PD-L1). 211 Negative controls were prepared by replacing the primary antibody with an irrelevant 212 one in all cases. 213

#### Immunohistochemical scoring

IHC stained slides were independently evaluated by 2 pathologists (L.M. and L.L.) 216 and discordant results were reviewed to reach a consensus. 217

TILs were evaluated on CD3- and CD20-stained slides and separately scored within 218 the tumor bulk (i.e. intratumor lymphocytes) and in the surrounding stroma (i.e. 219 peritumor lymphocytes). Briefly, intratumor lymphocytes were counted in 4 microscopic 220 fields at 200x and a score was arbitrarily assigned according to the following criteria: score 221 0 = < 10 positive lymphocytes; 1 = 10-25 positive lymphocytes; 2 = 26-60 positive 222 lymphocytes; 3 = > 60 positive lymphocytes. Peritumor lymphocytes were scored from 0-223 3 according to the number of lymphocytic aggregates in the tumor stroma, regardless of 224 their immunophenotype and counted in 10 fields at 4x magnification ("peritumor 225 lymphocyte total score"; score  $0 = \le 1$  aggregates; 1 = 2-6 aggregates; 2 = 7-14 aggregates; 226  $3 = \geq 15$  aggregates). In addition, the proportion of CD3 and CD20 positive lymphocytes 227 in peritumor aggregates was recorded. 228

Similarly, PD-1-expressing lymphocytes were separately scored in intratumor and 229 peritumor areas. For intratumor PD-1+ lymphocytes, the same scoring system described 230 above for CD3 and CD20 was applied. For peritumor lymphocytes, the percentage of PD- 231 1+ cells was assessed, and a semiquantitative score was assigned as follow:  $0 = \langle 5\%; 1 = 5 \rangle$  232 24%; 2 = 25-49%;  $3 = \geq 50\%$ . 233

For PD-L1 evaluation, the percentage and the intensity (0= negative; 1= mild; 2= 234 moderate; 3= marked) of positive tumor cells was assessed. 235

#### Statistical analysis

All statistical analyses were performed using R software. The association of clinico-238 pathological features with the IHC results was explored. Fisher's exact test was used for 239 categorical variables, while continuous variables were tested by means of Student t-test 240 or Wilcoxon rank-sum test, according to the results of Shapiro-Wilk test previously 241 performed to assess the normality of the distribution. The correlation between the IHC 242 results was investigated by Spearman rank correlation test. Survival analysis was 243 conducted using survival and survminer packages. Univariate Cox proportional-hazards 244 model was used to test the impact of the following variables on DFI and ST: age, sex, 245 breed, weight, presence of lymph nodes metastasis at the time of the diagnosis, presence 246 of hypercalcemia, tumor size (major diameter measured after formalin fixation), type of 247 treatment (adjuvant chemotherapy in addition to surgery), local recurrence, occurrence of 248 regional or distant metastasis, histopathological features and IHC results. A cut-off of 249  $p \le 0.05$  was used to screen the variables to be included in the multivariate analysis. Kaplan-250 Meier curves for categorical variables were drawn and compared by means of log-rank 251 test. 252

#### 3. Results

#### Study population

Forty-one dogs were included in the study (Table S1). Twenty-three (56.1%) were255purebred dogs, including Labrador retrievers (n=9), Border collies (n=3), Czechoslovakian256wolfdogs (n=3), German shepherds (n=2). Thirty-two (78%) dogs were female (5 intact; 27257spayed) and 9 (22%) were male (4 intact; 5 neutered). Median age at diagnosis was 11258years (range: 5-15 years) and median body weight was 24 kg (range: 7-40 kg).259

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Fifteen dogs (36.6%) were hypercalcemic at presentation and 3 (7.3%) had a bilateral260tumor mass. The median tumor size (major diameter measured after formalin fixation)261was 3.5 cm (range: 0.5-10 cm). Thirty-one (75.6%) dogs had sublumbar lymphadenopathy262at presentation, confirmed as metastatic AGASACA by histopathology in all cases.263Fourteen dogs (34.1%) were treated only by surgical resection of the primary tumor and264regional lymph nodes. Twenty-seven dogs (65.8%) received adjuvant chemotherapy,265which consisted of toceranib phosphate (n=25), carboplatin (n=1) and melphalan (n=1).266

#### Generation and validation of anti-PD-1 and anti-PD-L1 monoclonal antibodies

For PD-L1 antibodies, three clones were considered for further validation as they 270 detected PD-L1-FC recombinant protein in the dot blot. The clones were called 1.1 (IgG3), 271 2.1 (IgG2a) and 3.1 (IgM). All the three clones detected PD-L1-FC in ELISA (Figure S1 A-272 C) but none did in the FC assay against cells constitutively expressing PD-L1 (Figure S1 273 D-F, representing the 3.1 clone). In a WB against PD-L1-FC fusion protein, only one clone 274 (IgM PD-L1 3.1) reacted and a single band of approximately 56 kDa was detected (Figure 275 S2 A). Hence, only the IgM PD-L1 3.1 clone was used for further analysis and tested in 276 WB against the three canine cell lines (K9TCC, K9TCC-SH and DH82), producing a 277 dominating band of approximately 34 kDa and some weaker bands above it, likely 278 representing different post-translational forms of PD-L1 [34] (Figure S2 B). 279

For PD-1 antibodies, one clone, designated 1.1 (IgG2a), was chosen, which detected 280 rcPD-1-His in the dot blot. PD-1 1.1 detected PD-1-FC in ELISA in a dose-dependent 281 manner, within a range of 0.1-100 ng/well (Figure S3 A). When the antibody was tested by 282 flow cytometry against K9TCC and K9TCC SH cells, constitutively expressing canine PD-283 1, positive cells were identified (Figure S3 B, C). In addition, increased fluorescent signal 284 was detected in K9TCC cells overexpressing PD-1 (K9TCC-PD1 OE; Figure S3 D). In WB 285 the antibody produced a dominant band, approximately 70 kDa in size, with additional, 286 weaker bands underneath (Figure S4). Similarly to PD-L1, PD-1 was heavily glycosylated. 287 Interestingly, in DH82 cell line a single, weaker band at approximately 36 kDa was found, 288 which is close to the size of deglycosylated PD-1 (Figure S4). 289

#### Histopathology and immunohistochemistry

Histopathological features and IHC scores for each dog are reported in Table S1. TILs 292 were evaluated by IHC for CD3 (T lymphocytes) and CD20 (B lymphocytes), in addition 293 to the immune checkpoint molecule PD-1. 294

Intratumor lymphocytes were interspersed with tumor cells in the tumor bulk and 295 were predominantly CD3+ T cells, present in variable number: 2 cases scored 0; 15 cases 296 scored 1; 14 cases scored 2; 10 cases scored 3 (Figure 1A). CD20+ intratumor lymphocytes 297 were rarely detected (all cases scored 0) (Figure 1B). In 13 cases (31.7%), intratumor 298 lymphocytes also expressed PD-1 in a low percentage (score=1), and a positive correlation 299 between the number of CD3+ and PD-1+ intratumor lymphocytes was detected ( $\rho$ =0.36; 300 p=0.02) (Figure 1C). 301

Regarding peritumor lymphocytes total score, 1 case scored 0, 11 cases scored 1, 20 302 cases scored 2 and 9 cases scored 3. By IHC, peritumor lymphocytes were a mixture of 303 CD3+ and CD20+ cells, often distributed in nodular aggregates, composed of a variable 304 proportion of T and B lymphocytes (Figure 1D and 1E). Generally, CD3+ cells 305 predominated over CD20+ cells, with a CD3+/CD20+ cell ratio  $\geq 1$  in 98% (40/41) of the 306 cases. In 61% (25/41) of the cases, a variable proportion of peritumor lymphocytes also 307 expressed PD-1 (11 with score 2; 14 with score 3) (Figure 1F). Local recurrence was 308 significantly associated with a low peritumor lymphocyte total score (p=0.009) and a low 309 CD3+/CD20+ ratio in peritumor lymphocytes (p=0.034). 310

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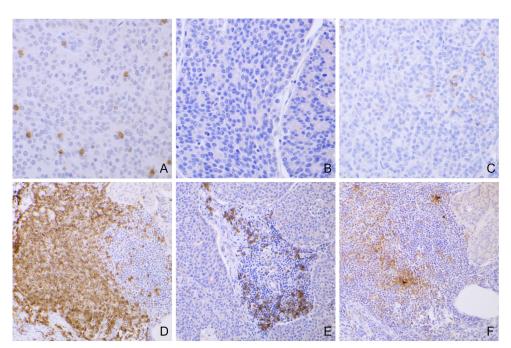


Figure 1. Tumor infiltrating lymphocytes in canine AGASACAs. (A) Numerous 313 intratumor CD3+ lymphocytes (case #22; 40x magnification). (B) Absence of intratumor 314 CD20+ lymphocytes (case #22; 40x magnification). (C) Scattered intratumor PD-1+ 315 lymphocytes (case #22; 40x magnification). (D, E, F) Peritumor lymphocytes in nodular 316 aggregates composed of a mixture of CD3 (D), CD20 (E) and PD-1 (F) positive 317 lymphocytes (case #5; 20x magnification). 318

Seventeen AGASACAs (42%) expressed PD-L1 in a range between 5% and 95% of 321 tumor cells, with mild to moderate intensity (Figure 2). Interestingly, a single case (case #36) was characterized by a regional area of undifferentiated tumor cells, displaying highly infiltrative behavior, spindeloid morphology and increased cellular atypia. These cells had diffuse and intense expression of PD-L1. No associations between PD-L1 325 expression and other IHC scores were identified.

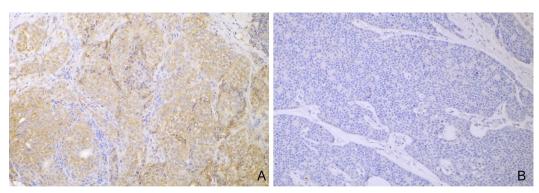


Figure 2. PD-L1 expression in canine AGASACAs (20x magnification). (A) 329 Representative case with diffuse and moderate expression of PD-L1 (case #18). (B) Negative case (case #20).

Survival analysis

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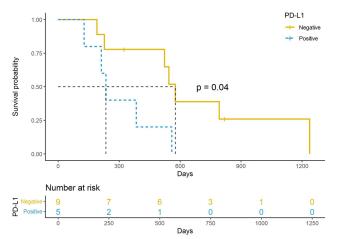
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At the time of data analysis, 5 dogs (12.2%) were alive after 39 to 819 days, 22 (53.7%) 335 had died for causes attributable to AGASACA (range: 128-1237 days), 14 (34.1%) for 336 unrelated causes (range: 108-1434 days) and one was lost to follow-up (598 days). Eight 337 dogs (19.5%) experienced metastatic disease to regional (ileosacral) lymph nodes after the 338 primary surgery, while 11 dogs (26.8%) developed distant metastasis to various organs, 339 including distant lymph nodes, spleen, liver, and lungs. The median DFI of the whole 340 population was 474 days (range: 0-1237 days), and median ST was 472 days (range: 108-341 1434 days). By univariate Cox proportional-hazards analysis, dogs with metastatic 342 regional lymph nodes at presentation had a significantly shorter DFI (369 vs 940 days; 343 p=0.046) (Table S2). Body weight, local recurrence, and histological presence of 344 lymphovascular invasion and tumor necrosis negatively affected the ST, however only 345 body weight and lymphovascular invasion maintained statistical significance in 346 multivariate analysis (p=0.027 and p=0.012, respectively) (Table S2). 347

TILs and immune checkpoint molecules expression did not influence the outcome in348the whole population. When the subgroup of dogs treated by surgery alone (n=14) was349considered, tumor expression of PD-L1 was associated with a significantly shorter ST350compared to cases that tested negative for the marker (576 vs 235 days; p=0.022) (Figure351352352



*Figure 3.* Kaplan–Meier curves of survival time of 14 dogs with AGASACA treated 353 with surgery alone. 354

4. Discussion

Deciphering the interaction between the tumor and the patient's own immune 358 system has led to the development of cancer immunotherapies, which recently 359 revolutionized the paradigm of cancer treatment in human oncology. In this context, 360 adaptative immunity, and intratumor lymphocytes in particular, was shown to play a key 361 role in the antitumor immune response [35]. Based on the degree of lymphocytic 362 infiltration and molecular signatures, tumors can be classified in three different 363 phenotypes (inflamed, immune-excluded, immune-desert), that are strongly associated to 364 the response to immunotherapeutic treatments, with inflamed (hot) tumors being more 365 likely to benefit from the administration of checkpoint inhibitors [36]. Given the 366 recognized role of several canine tumor types as a spontaneous model for human 367 oncology, there is an increasing interest in characterizing the immune landscape of these 368 tumors. Osteosarcoma, oral malignant melanoma, lymphoma, and mammary carcinomas 369 belong to canine tumors presenting a hot immune phenotype [22,37]. 370

To the best of our knowledge, the immune microenvironment of AGASACA has not 371 been fully characterized so far, but a recent study suggested a possible association, even 372

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if not statistically significant, between inflammatory cell infiltration and outcome [19]. In addition, transcriptome analysis confirmed a "hot" immune signature [38]. 374

Here, in order to further characterize the AGASACA at the molecular level, we have 375 developed two new antibodies against canine ICs highly relevant to cancer in both 376 humans and dogs: PD-1 and PD-L1. Preliminary validation revealed that the PD-1 377 monoclonal antibody detects its target in WB, ELISA, and FC experiments. Among the 378 three developed PD-L1 clones, all were effective in ELISA, none in FC, and only the 3.1 379 clone in WB. Upon basic validation, the antibodies PD-L1 3.1 and PD-1 1.1 were used to 380 assess target expression on the canine AGASACA samples. The antibodies will be further 381 characterized in future to assess their potential as drug candidates. 382

In the present study, the presence and phenotype of TILs was evaluated by IHC in 383 41 surgically resected AGASACAs and a high degree of lymphocytic infiltration, both in 384 peritumoral and intratumor location, was observed. Moreover, while peritumor 385 lymphocytes were a mixture of T- and B-cells, intratumor lymphocytes were almost 386 exclusively represented by CD3+ T-cells. These findings support an inflamed 387 immunophenotype at least for a subset of cases; in fact, contrary to the immune-excluded 388 phenotype, which is characterized by lymphocytic infiltrates limited to the tumor stroma, 389 inflamed tumors are defined by the presence of T-cells intermingled with neoplastic cells 390 [36]. Even if molecular investigations were beyond the aim of our study, the 391 immunohistochemical characterization of TILs suggests a "hot" phenotype of canine 392 AGASACAs and encourage further investigations in this direction. 393

Cancer immunotherapy in veterinary oncology is still at early stages of development 394 and application. Some successful examples are represented by the clinical implementation 395 of autologous vaccines in canine lymphoma [39] and anti-CSPG4 electro-vaccination in 396 canine melanoma [40]. Another early but promising example is represented by the empty 397 cowpea mosaic virus treatment in inflammatory mammary carcinoma [41]. Only few 398 preliminary studies, by using specific canine anti-PD-L1 and anti PD-1 antibodies for the 399 treatment of oral malignant melanoma, demonstrated the potential value of immune 400 checkpoint blockade in dogs [24,26]. To evaluate if canine AGASACA is a possible 401 candidate for immunotherapy, the study population was tested by IHC with monoclonal 402 antibodies specifically targeting canine PD-1 and PD-L1, and a subset of TILs expressed 403 PD-1 in variable proportions. These were localized both within the tumor bulk (i.e., 404 intratumor) and in the surrounding stroma (i.e., peritumor); moreover, in intratumor 405 areas the number of PD-1+ lymphocytes positively correlated with the number of CD3+ 406 cells. Although no prognostic value of PD-1 expression by TILs was demonstrated, the 407 enrichment of the tumor microenvironment with PD-1-positive lymphocytes might 408 support their role as a possible target for the development of immunotherapeutic 409 strategies. 410

To date, the expression of PD-L1 at the protein level has been investigated in a variety 411 of canine tumors. The most comprehensive studies, which included preliminary clinical 412 trials or PD-L1 association with clinico-pathological features and/or outcome, focused on 413 melanoma, osteosarcoma, and mammary carcinoma [21,25,26,42]. In addition, two studies 414[26,43] tested the expression of PD-L1 by IHC in a limited number but wide range of 415 canine tumors, demonstrating its expression in several tumor types, including 19 out of 416 20 of the tested AGASACAs. Among these, most of the tumors expressed PD-L1 in more 417 than 50% of the tumor cells. Similarly, in the present study, the expression of PD-L1 by 17 418 out of 41 AGASACAs was demonstrated, with variable percentages of positive tumor 419 cells. Lack of standardized methods to assess PD-L1 expression by tumor cells in 420 veterinary medicine might have contributed to the lower proportion of positive 421 AGASACAs observed in our study compared to the previous one [26]. In this context, it 422 is also worth to mention the variability observed in human oncology, when different 423 antibody clones and staining platforms are used to assess PD-L1 tumor proportional score 424 [44]. Interestingly, a single case in this study had a regional tumor area displaying 425 morphological features of undifferentiated carcinoma with highly invasive growth 426 pattern; this area was also characterized by diffuse and intense positivity for PD-L1. Given427the morphological features and the early recurrence and regional metastasis observed in428this dog, this finding might suggest an association between PD-L1 activation and tumor429aggressive behavior.430

Survival analysis for the whole cohort showed that body weight, local recurrence, 431 and histological presence of lymphovascular invasion and tumor necrosis negatively 432 affected the outcome, as previously described [18,19]. A shorter DFI was also observed in 433 dogs with nodal metastasis at presentation. 434

PD-L1 expression was associated with a poorer outcome only in the subgroup of dogs 435 treated with surgery alone (i.e., without medical treatment). This result indicates that, 436 similarly to other canine tumors [21,23] PD-L1 expression in AGASACAs influenced 437 tumor behavior, affecting the ST, thus representing a potential prognostic biomarker. 438 However, due to the limited patient number, this preliminary result needs to be 439 interpreted with the consideration of potential bias and limitations. Future studies, on a 440larger cohort of patients to confirm this result are needed. The lack of association between 441 PD-L1 expression and survival when the whole cohort was considered might be explained 442 by the influence that adjuvant treatment with toceranib phosphate has on survival 443 [4,12,13]. In our study, we did not observe a clear impact of chemotherapy treatment on 444 survival, but since the chemotherapy protocol was not standardized, it is possible that the 445 adjuvant treatment might have represented a confounding factor when the whole 446 population was considered, potentially obscuring the impact of other variables on the 447 outcome. To our knowledge, the direct effect of toceranib phosphate on PD-L1 expression 448 in canine cancer has not been investigated so far . However, there is some evidence for 449 induction of PD-L1 destabilization and degradation by sunitinib, a tyrosine kinase 450 inhibitor similar to toceranib, leading to a decreased expression of this molecule on tumor 451 cells [45]. Although not yet confirmed, the effect of toceranib might be similar. 452

#### 5. Conclusions

In conclusion, our study showed a high and heterogenous expression of immune 454 checkpoint molecules and the presence of a lymphocytic infiltration in canine 455 AGASACAs. PD-L1 was also associated with the outcome in a subset of patients, 456 indicating a prognostic significance for this marker. Even if in our cohort TILs were not 457 associated with the outcome, the high level of lymphocytic infiltration suggests a hot 458 immune phenotype of this tumor. Taken together, these results show the relevance of the 459 immune-microenvironment in canine AGASACAs, indicating that the stimulation of anti-460 cancer immunity might represent a valuable treatment approach for this tumor type and 461 deserve more in-depth investigations. 462

Supplementary Materials: The following supporting information can be downloaded at:463www.mdpi.com/xxx/s1, Figure S1: PD-L1 antibody validation: ELISA and FC; Figure S2: PD-L1464antibody validation: Western Blot; Figure S3: PD-1 antibody validation: ELISA and FC; Figure S4:465PD-1 antibody validation: Western Blot; Table S1: Clinicopathological features; Table S2: Survival.466

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