Identification of CSPG4 as a promising target for translational combinatorial approaches in osteosarcoma

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Abstract

Background: Osteosarcoma (OSA) is a highly metastatic pediatric bone tumor. Adjuvant chemotherapy and surgical resection represent standard treatments; however, the prognosis is still poor. Effective strategies are urgently needed. Chondroitin sulfate proteoglycan (CSPG)4 is a transmembrane proteoglycan with a low expression in normal tissues but high expression in several solid tumors, where it plays a central tumorigenic role. Therefore, it represents a promising therapeutic target. The high homology between human and canine CSPG4 and the recognized translational power of canine tumors as preclinical models for human malignancies prompted us to evaluate CSPG4 expression and the consequences of its immune-targeting for both human and canine OSA treatment.

Methods: We analyzed CSPG4 overexpression in human and canine OSA samples and its significance for the survival of OSA patients. We exploited functional in vitro experiments to assess the antitumor potential of CSPG4 immune-targeting.

Results: CSPG4 is overexpressed in OSA and has possible clinical implications as suggested by an evident correlation between CSPG4 overexpression and a shorter survival for both OSA-affected humans and dogs. The potential of CSPG4 immune-targeting for OSA treatment came from the ability of anti-CSPG4 monoclonal antibodies and sera, derived from human-CSPG4-DNA vaccinated canine patients, to significantly inhibit human and canine CSPG4-positive OSA cell proliferation, migration, and osteospheres generation. Moreover, CSPG4 immune-targeting has been shown to potentiate the effect of doxorubicin.

Conclusions: Overall, these results provide the rationale to investigate the CSPG4 immune-targeting as a promising weapon for the treatment of CSPG4-positive OSA canine patients, to be successfully translated to a human setting.

Keywords: antibodies, comparative oncology, CSPG4, osteosarcoma

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Introduction

Osteosarcoma (OSA) is the most common primary malignant tumor of the bones. Of note, OSA is the sixth most-frequent pediatric cancer and represents the second most common cause of cancer-related death in this age group; a second peak is observed in adults after the sixth decade of life.1,2 Currently, primary OSA is classified into low- (Grade I) and high-grade subtypes (Grade II and III in the presence of metastasis), high-grade OSA being the most prevalent and aggressive variant.3

At initial diagnosis, almost 20% of OSA patients present evidence of metastatic spreading commonly involving lungs (90%), sites within the same affected bones (8–10%) or, more rarely, lymph nodes.4,5 However, considering that the
vast majority of patients without overt metastasis at diagnosis develop lung metastasis within 6–36 months, it is presumed that these apparently non-metastatic patients actually have micrometastases already at diagnosis.6,7

Conventional treatments consist mainly in the surgical resection of the primary tumor, in association with neoadjuvant and adjuvant chemotherapy with doxorubicin, methotrexate, and cisplatin. On the basis of the percentage of tumor necrosis after neoadjuvant chemotherapy, patients can be classified as poor responders (Huvos Grades I–II), responders (Huvos Grade III), or good responders (Huvos Grade IV). A strong correlation between the Huvos Grade and the subsequent effectiveness of postoperative chemotherapy and disease-free survival has been observed. This standard of care treatment is quite effective in the setting of localized OSA. Indeed, nowadays, the 5-year survival rate for localized OSA is about 65–70%, while only 20% of patients with multifocal disease or metastasis at diagnosis are still alive after 5 years.2,7,8

Therefore, it is evident that OSA is still a very aggressive and fatal disease for which no significant therapeutic advances have been achieved in the last 30 years and for which the identification of novel and more effective approaches is needed to improve patient survival.

As a step forward, during the last few decades, researchers have addressed to the identification of potentially therapeutically targetable OSA-associated antigens. Several tyrosine kinase receptors have been identified as over-expressed in OSA, including KIT, vascular endothelial growth factor receptor (VEGFR)-2 and VEGFR-3, platelet-derived growth factor (PDGFR)-β, and MET, and found to be correlated with metastasis development and poor survival.9–12 As a consequence, in the last few years several targeted therapies have been investigated, which unfortunately showed only limited efficacy in advanced OSA.11,13 The need for novel and more clinically relevant targets is therefore critically evident.

The chondroitin sulfate proteoglycan (CSPG)4 is a cell surface proteoglycan, considered as an ideal tumor-associated antigen, that is, an oncoantigen,14 because it is poorly expressed in healthy tissues,15–23 whereas in a vast range of human neoplasms it is in overexpressed by tumor cells, the tumor microenvironment and, cancer stem cells (CSCs). It has been widely described that CSPG4 has a key role in several oncogenic pathways required for malignant progression and metastasization.24 We have already demonstrated the clinical relevance of CSPG4 immune-targeting by means of DNA vaccination for the treatment of canine malignant melanoma (MM),25,26 however, to the best of the authors’ knowledge very few investigations have been undertaken on the involvement of CSPG4 in OSA.27,28

The aim of this study was to evaluate CSPG4 as a potential immunotherapeutic target for both human and canine OSA patients. This could offer an appealing opportunity to exploit spontaneous occurring OSA in pet dogs as model of human OSA tumors and for predicting the clinical efficacy of therapeutic approaches targeting CSPG4. Indeed, canine OSA is nearly indistinguishable from the human disease, presenting the same risk factors, histological tumor grading, similar standard treatments, and clinical responses.29,30 All these shared features make OSA-bearing dogs a valuable translational model of human OSA for the investigations of novel immunotherapies that could benefit both species.

Based on these considerations, we investigated CSPG4 expression in both human and canine OSA. We aimed to evaluate the effects of anti-CSPG4 targeting, alone or in combination with doxorubicin, on both human and canine OSA tumor cells and on osteospheres, enriched in CSC, which are considered responsible for recurrences and metastasis. Overall, our results provide the rationale for testing the clinical effectiveness of an anti-CSPG4 immunotherapy in dogs affected by spontaneous OSA, with the final aim of its translation for the treatment of human OSA patients.

Material and methods

Sample collection and clinical follow up

Tissue samples from 50 cases of spontaneous canine appendicular OSA collected via routine care between 2008 and 2014 at the Veterinary Teaching Hospital, Department of Veterinary Science of the University of Turin, were examined in this study.

Client-owned dogs affected by spontaneous OSA were treated according to the European guidelines established in the Principles of Laboratory
Animal Care (directive 86/609/EEC). The Ethical Committee of the Department of Veterinary Science (University of Turin) approved the experimental protocol, which follows the best practice of veterinary care; written consent for entry into the study was obtained from dog owners. For all canine OSA patients, the initial data collected included history, physical examination, complete blood count, serum biochemical profile, urinalysis and abdominal ultrasound. Limb [lateral–lateral (LL) and anterior–posterior (AP) views] and chest [LL, right and left, and dorso–ventral (DV) views] radiographic evaluation was performed to examine the features and the extent of the tumor in addition to the presence of lung metastasis. From 2010 evaluation with computer tomography (CT) was included. In cases where regional lymph nodes were enlarged, they were aspirated, cytologically examined, and then removed for histological evaluation. All dogs included in this study were surgically treated (amputation or limb sparing) before receiving adjuvant chemotherapy. The Kaplan–Meier method was used to estimate overall and disease-free survival times. Differences in survivals were tested for significance using log-rank tests.

**Immunohistochemical analysis**

Immunohistochemical (IHC) analysis was performed as described previously on collected canine OSA. Samples were fixed in 4% neutral buffered formalin, embedded in paraffin, and sectioned at 4 μm. Then, samples were stained with hematoxylin and eosin to establish the histological diagnosis. The histological grade was determined according to the systems proposed by Loukopoulos and Robinson. The grading was evaluated as I (low), II, or III (high). IHC for CSPG4 in the 50 samples was performed on 4 μm sections of formalin-fixed, paraffin-embedded tissues. Sections were exposed to high-temperature for antigen unmasking by incubation at 98°C with citric acid buffer, pH 6.0. Endogenous peroxidase activity was blocked with 3% hydrogen peroxide in methanol for 30 min at room temperature. Tissue sections were incubated for 12 h at room temperature with a polyclonal anti-CSPG4 antibody (diluted 1:40, Sigma Aldrich, St. Louis, MO, USA), then 30 min with biotinylated-secondary antibody (Vectastain Elite ABC) and revealed with the ImmPACT DAB kit for peroxidase, both from Vector Laboratories Inc. (Burlingame, CA). A total score considering the proportion of positively stained tumor cells and the average staining intensity was assigned as previously described. Briefly, the score indicating the positivity of tumor cells was assigned as follows: 0 (none); 1 (<1/100 or <1%); 2 (1/100–1/10 or 1–10%); 3 (1/10–1/3 or 10–30%); 4 (1/3–2/3 or 30–70%); and 5 (>2/3 or >70%). The score representing the estimated average staining intensity of positive tumor cells encompass 0 if none, 1 weak, 2 intermediate, or 3 strong. The two scores were then added to each other to obtain a final score of CSPG4 expression ranging from 2 to 8. A total score ≥4 was used as cut-off.

**Cell lines and osteospheres**

Human OSA cell lines (MG-63, SAOS-2, U2OS) were obtained from the American Type Culture Collection (ATCC, Manassas, VA, USA) and were cultured in Dulbecco’s Modified Eagle Medium (DMEM; Sigma Aldrich) or RPMI (Sigma Aldrich) supplemented with 20% fetal bovine serum (FBS, Sigma Aldrich). Penny cells, derived from the tail biopsy of a primary grade III canine OSA, were grown in ISCOVE Modified Dulbecco’s Medium (Sigma Aldrich) supplemented with 10% FBS. All cells were grown in medium supplemented with penicillin/streptomycin (Sigma Aldrich) and maintained at 37°C and 5% carbon dioxide in a humidified incubator.

Both human and canine osteospheres were generated following the protocol described in Conti et al. Briefly, cells were detached and plated in ultra-low-attachment 75 cm flasks (Sigma Aldrich) at 6 × 10^4 viable cells/ml in serum-free DMEM-F12 (Sigma Aldrich) supplemented with 0.4% bovine serum albumin (BSA), 20 ng/ml basic fibroblast growth factor (bFGF), 20 ng/ml epidermal growth factor (EGF), 5 mg/ml insulin, all from Sigma Aldrich. Non-adherent spherical cells’ clusters, named osteospheres, were collected after 48 h or 5 days for further analysis. Photographs of osteospheres were taken using a CCD-300-RC camera, and images were processed using Fiji Software (Rasband, W.S., ImageJ, US National Institutes of Health, Bethesda, MD) and PowerPoint (Microsoft, Redmond, WA).

**MTT cell proliferation assay**

3-(4,5-dimetiltiazol-2-il)-2,5-difeniltetrazol (MTT; Merck Millipore, Burlington, MA, USA)
assay was used to assess proliferation of human MG-63 and canine Penny OSA cells. Briefly, epithelial cells were seeded in triplicate in 96-well plates \((5 \times 10^3\) cells and \(8 \times 10^3\) cells/100µl well, respectively) in serum-free medium and allowed to adhere overnight. To evaluate the inhibition of the CSPG4-mediated cell proliferation, four selected anti-CSPG4 purified mAbs, 225.28, TP32, TP49, and VF20-VT87.41, were mixed in a pool or used as single agents. All the mAbs were produced in the laboratory of Prof. Ferrone (Massachusetts General Hospital, Harvard Medical School, Boston, MA) and are secreted by hybridomas generated from BALB/c mice immunized with human melanoma cells. Specifically, the mAb 225.28 is a mouse IgG2a, while the others (TP32, TP49, and VF20-VT87.41) are all mouse IgG1, which recognize distinct and spatially distant epitopes of human CSPG4. The specificity of mAbs was characterized as described elsewhere.\(^{34-36}\)

Cells were incubated with different treatments for 48h at the following final concentrations or dilutions: control isotypes (Sigma Aldrich; 25µg/ml), anti-CSPG4 mAbs (225.28, TP32, TP49, and VF20-VT87.41), tested individually at 25µg/ml or mixed in a pool to a final total concentration of 25µg/ml, doxorubicin (Sigma Aldrich; 100 nM, 1µM, or 10µM), melanoma canine patients’ sera collected pre- and post-vaccination (1:100) with Hu-CSPG4 DNA plasmid.\(^{25,26}\) Cells grown in medium alone were used as control. MTT solution (5mg/ml) was added to each well following manufacturer’s instruction. Following 4h incubation at 37°C, 100µl of dimethyl sulfoxide (DMSO, Sigma Aldrich) were added to dissolve formazan crystals and absorbance was measured on an enzyme-linked immunosorbent assay (ELISA) plate reader (Bio-Rad, Hercules, CA) with a wavelength of 570 nm.

To assess viability of human MG-63 and canine Penny osteosphere-derived cells, \(6 \times 10^4\) cells/ml were seeded in 96-well plates in non-adherent conditions and incubated with different stimuli as described above and spheres were allowed to generate for 48h. Following overnight incubation at 37°C, formazan crystals were dissolved by adding 100µl isopropanol with HCl 0.04N to each well. Optical density was measured on an ELISA plate reader (Bio-Rad) with a test wavelength of 570 nm and a reference wavelength of 630 nm. Difference between 570 nm and 630 nm readings represents the output value.

**Flow cytometry analysis**

Human and canine OSA epithelial cells and osteosphere were detached and dissociated by using Cell Dissociation Non Enzymatic 1X solution (Sigma Aldrich) and then resuspended in the culture medium with a concentration of \(1 \times 10^5\) cells in 100µl. Cytofluorimetric analysis were performed on cells for the detection of CSPG4 antigen using a mixed pool of the following mAbs (225.28, TP32, TP49, and VF20-VT87.41; 25µg/ml final concentration) directed towards different epitopes of CSPG4 and produced in the lab of Prof. Ferrone (refer to MTT cell proliferation assay). Pooled mAbs were incubated on OSA cells for 30min at 4°C and antibody binding was evaluated using a fluorescein isothiocyanate (FITC)-conjugated rabbit anti-mouse Ig (Dako-Cytomation, Glostrup, Denmark). Flow cytometry was performed with a FACS Verse (BD Biosciences, San Jose, CA, USA) and the results were expressed as percentage of positive cells and mean fluorescence intensity (MFI) and analyzed with BD Facs Suite software.

**Western blotting**

Human and canine OSA cells were incubated in lysis buffer (150mM NaCl, 50mM Tris-HCl, pH 8.0, 1% Nonidet P-40, 1mM Na3VO4, 1mM NaF, and protease inhibitors, all from Sigma Aldrich) for 30min at 4°C. Cell lysates were centrifuged for 5min at 15,000g. Total protein concentration was quantified using the Pierce™ BCA Protein Assay Kit (Thermo-Fisher Scientific, Rockford, IL, USA). Following 5’ denaturation at 95°C in 2-mercaptoethanol-containing Laemmli sample buffer (Bio-Rad, Hercules, CA), equal amounts of protein (70µg) were separated through electrophoresis in an Any kDa Mini-Protein TGX precast gel (Bio-Rad) and then transferred onto an Immobilion-P PVDF membrane (Merck Millipore, Billerica, MA). Following blocking with 5% nonfat dry milk (Santa Cruz Biotechnology, Dallas, TX) in wash buffer (Tris Buffered Saline - TBS - supplemented with 0.05% Tween-20 from Sigma Aldrich), the membrane was incubated overnight at 4°C with a pool of anti-CSPG4 mAbs (225.28, TP32, TP49, and VF20-VT87.41, 5µg/ml each), washed three times in TBS and 0.05% Tween-20, and then incubated for 1h at room temperature with HRP-conjugated rabbit antimouse Ig (Sigma Aldrich). Actin was used as control for equal protein loading. Immunoreacting bands were detected using ECL (Thermo Scientific) according to the manufacturer’s instructions. Band relative intensity
was acquired using a ChemiDoc™ Touch Imaging System (Bio-Rad).

Cell migration assay
To measure cell migration, human MG-63 and canine Penny OSA cells were pre-incubated with different treatments at the following final concentrations: control isotypes (Sigma Aldrich; 100 µg/ml), anti-CSPG4 mAbs (225.28, TP32, TP49, and VF20-VT87.41 mixed in a pool to a final total concentration of 100 µg/ml), alone or in combination with doxorubicin (Sigma Aldrich; 10 µM or 100 nM); then seeded (5 × 10⁴ and 3 × 10⁴, respectively, per well) in 100 µl of serum-free medium in the top chamber of a 24-Transwell plate (8-µm pore size; Corning, Amsterdam, Netherlands). Cells incubated with medium alone were used as control. All bottom chambers of the Transwell plates were filled with 10% FBS-supplemented medium (600 µl per well) and cells were incubated at 37°C in a 5% CO₂ atmosphere. After 48h, the nonmigrated cells on the top side of the filter were removed by scrubbing twice with cotton tipped swab. Migrated cells on the bottom side of the filter were fixed with 2.5% glutaraldehyde (Sigma-Aldrich) and stained with 0.2% crystal violet (Sigma-Aldrich). After washing, the migrated cells of four randomly selected fields per well were imaged using an Olympus BX41 microscope (Olympus Corp., Tokyo, Japan) and analyzed using Fiji and ImageJ Softwares (Rasband, W.S., ImageJ, US National Institutes of Health).

Dataset
Genome-wide gene expression analysis was based on previously deposited microarray experiments and data preprocessing, described in Kuijjer et al. The dataset included genome-wide gene expression data of osteoblasts (GEO accession number GSE33382), mesenchymal stem cells (MSCs; GEO accession number GSE28974), and 84 high-grade OSA pre-treatment biopsies (GEO accession number GSE33382). CSPG4 mRNA expression resulted significantly upregulated in MSC, considered potential precursors for OSA development, as compared with normal osteoblasts. CSPG4 mRNA expression was further significantly increased in high-grade OSA biopsies (Figure 1(a)). Taking advantage of the clinicopathological details regarding the high-grade OSA samples included in the analyzed dataset, we sought to evaluate whether the CSPG4 expression level in pretreatment OSA biopsies could be related to their response to neoadjuvant chemotherapy, expressed in Huvos Grades. As shown in Figure 1(b), CSPG4 mRNA expression in pretreatment OSA samples is not related to the Huvos Grade, therefore it cannot be considered as a prognostic factor implicated in the response to the neoadjuvant treatment. Nevertheless, CSPG4 mRNA expression is quite higher in Huvos Grade I–III samples, which include OSA with a lower necrosis in response to neoadjuvant chemotherapy and therefore with a worse prognosis, as compared with Huvos Grade

Results
CSPG4 is overexpressed in human high-grade OSA and is related to a poor prognosis.

In order to evaluate the potentiality of CSPG4 as a novel target for human OSA, we used a publicly available comprehensive microarray dataset (super-series accession number GSE42352), including normalized gene expression data of osteoblasts (n = 3; GEO accession number GSE33382), MSCs (n = 12, GEO accession number GSE28974), and high-grade OSA pretreatment biopsies (n = 84, GEO accession number GSE33382). CSPG4 mRNA expression resulted significantly upregulated in MSC, considered potential precursors for OSA development, as compared with normal osteoblasts. CSPG4 mRNA expression was further significantly increased in high-grade OSA biopsies (Figure 1(a)). Taking advantage of the clinicopathological details regarding the high-grade OSA samples included in the analyzed dataset, we sought to evaluate whether the CSPG4 expression level in pretreatment OSA biopsies could be related to their response to neoadjuvant chemotherapy, expressed in Huvos Grades. As shown in Figure 1(b), CSPG4 mRNA in pretreatment OSA samples is not related to the Huvos Grade, therefore it cannot be considered as a prognostic factor implicated in the response to the neoadjuvant treatment. Nevertheless, CSPG4 mRNA expression is quite higher in Huvos Grade I–III samples, which include OSA with a lower necrosis in response to neoadjuvant chemotherapy and therefore with a worse prognosis, as compared with Huvos Grade
IV samples, characterized by OSA with a 100% necrosis after neoadjuvant chemotherapy and with a better prognosis. In addition, CSPG4 overexpression showed a trend of correlation with a shorter overall survival (OS; Figure 1(c)) and metastasis-free survival (Figure 1(d)) of OSA patients. Taken together, these data suggest the potential role of CSPG4 in human OSA progression and the possible clinical relevance of its adjuvant targeting for high-grade OSA treatment, representing an even more interesting opportunity for those OSA that are not responsive to neoadjuvant chemotherapy.

CSPG4 is highly expressed in canine OSA
Canine patients affected by spontaneous OSA are considered a highly translational and predictive model to evaluate the clinical response to innovative therapeutic treatments for human OSA, providing several advantages over mouse models. On the basis of these considerations and to confirm the predictivity of the canine OSA model also for CSPG4, first we investigated its expression in canine OSA biopsies. Specifically, we evaluated CSPG4 protein expression using IHC in a total of 50 samples of canine appendicular OSA from patients treated between 2008 and 2014 at the Department of Veterinary Science of the University of Turin. IHC staining for CSPG4 detection was performed on all the collected tissues as previously described in order to obtain a value of 0 (if negative) or from 2 to 8. There were 38 (76%) CSPG4-positive and 12 (24%) CSPG4-negative primary canine OSA samples (Figure 2(a)). For all CSPG4-positive samples, staining was observed mainly on the membrane surface and in the
Figure 2. Expression of chondroitin sulfate proteoglycan (CSPG4) antigen in canine osteosarcoma (OSA) samples. (a) Percentage of primary OSA that scored positive (black bar) or negative (gray bar) for CSPG4 expression. (b) Immunohistochemical (IHC) staining of canine OSA biopsies with an anti-CSPG4 antibody. Representative sections of primary OSA not expressing (1), expressing low levels (2), or expressing high levels (3–6) of the CSPG4 antigen. (c) Percentage of CSPG4-positive primary OSA considering the different histological grade. (d) CSPG4 score of expression (mean ± SD) of CSPG4-positive primary OSA considering the different histological grade.
cytoplasm of neoplastic cells, while no staining was detected in the surrounding healthy tissue (Figure 2(b)). The score of CSPG4 expression was different among positive samples and representative IHC examples are shown in Figure 2(b). Considering the histological classification of OSA biopsies at diagnosis, CSPG4 expression was observed in 10 out of 12 (83%) canine patients of Grade I, 17 out of 21 (81%) of Grade II, and 11 out of 17 (65%) of Grade III (Figure 2(c)). CSPG4 expression score showed to be independent from the tumor grade (Figure 2(d)). Overall, these results labeled the CSPG4 as a tumor-associated antigen overexpressed in canine OSA.

CSPG4 overexpression in canine OSA is related to poor survival

For each of the 50 canine patients included in the study, in addition to the OSA biopsy, we collected clinical data at the diagnosis including history, a physical exam, complete blood count, serum biochemical profile, urinalysis, abdominal ultrasound, and clinical follow up. Limb and chest radiographic (or CT) evaluation was performed to examine the features and the extent of the tumor and to exclude lung metastasis at diagnosis. All dogs included in this study were surgically treated (amputation or limb sparing) before receiving adjuvant chemotherapy using doxorubicin (30 mg/m², 4–5 administrations, 21 days apart) or cisplatin (70 mg/m², 4–5 administrations, 21 days apart) as single agents or in combination (4 cycles, 21 days apart, each cycle consisting of cisplatin 50 mg/m² at day 1 and doxorubicin 15 mg/m² at day 2).

Canine patients were clinically and radiographically examined every 3 months during the first year after the conclusion of chemotherapy and then every 6 months. OS was considered as the days between the surgery and death, while the disease-free interval (DFI) was considered as the number of days between surgery and tumor recurrence and/or evidence of metastatic disease. Considering a CSPG4 score $\geq$ 4 as threshold, canine patients affected by CSPG4-positive OSA displayed a significantly shorter survival as compared with CSPG4-negative OSA (Figure 3(a)), with a median survival time (MST) of 271 and 440 days, respectively (Table 1). As far as the DFI is concerned, a trend is evident (Figure 3(b)), with a median DFI of 237 days for canine patients bearing a CSPG4-positive OSA and 339 days for dogs affected by a CSPG4-negative OSA (Table 1).

Overall, these results suggest the potential clinical relevance of CSPG4 for canine OSA progression, highlighting the importance of its targeting for comparative oncology studies for OSA treatment.

CSPG4 immune-targeting significantly inhibits CSPG4-dependent human and canine OSA cells proliferation

As CSPG4 facilitates constitutive activation of signaling pathways, which promote cell proliferation, we sought to in vitro evaluate the impact of CSPG4 immune-targeting on the growth of OSA cells. First, we evaluated CSPG4 expression in human (MG-63, U2OS, and SAOS-2) and canine (Penny) OSA cell lines. Flow cytometry analyses were performed using a pool of specific anti-CSPG4 mAbs (225.28, TP32, TP49, andVF20-VT87.41). Interestingly, cytofluorimetric results demonstrated that all the tested cell lines expressed high levels of CSPG4 (Supplemental Figure S1(a)) and this was confirmed also by Western blot analysis (Supplemental Figure S1(b)). For the following experiments, we selected the human MG-63 and the canine Penny cell lines. To understand the role of CSPG4 in OSA growth, MG-63 and Penny cells were incubated with control isotypes or anti-CSPG4 mAbs (225.28, TP32, TP49, andVF20-VT87.41), used as a mixed pool or as single agents. The proliferation of both OSA cell lines was significantly inhibited, as compared with control, by anti-CSPG4 mAbs used as a pool (Figure 4(a) and (b)) or as single agents (Supplemental Figure S2(a) and (b)). For this reason, in the following studies the mixed pool of the four anti-CSPG4 mAbs was used. As doxorubicin is a chemotherapeutic drug commonly used for the treatment of both human and canine OSA, we evaluated whether the CSPG4 targeting through mAbs could enhance its antiproliferative effect. At the lowest selected dose of doxorubicin (100 nM), MG-63 cells resulted resistant to the chemotherapeutic agent (Figure 4(a)), therefore the reduction in cell proliferation obtained by the combination of doxorubicin and anti-CSPG4 mAbs was primarily due to CSPG4 immune-targeting alone. As expected, when resistance is overcome using a higher dose of doxorubicin (10 µM), the MG-63 cell proliferation is significantly inhibited (Figure 4(a)). In this case, the combination with anti-CSPG4 mAbs is more effective in inhibiting human OSA cell proliferation compared with both the highest dose of doxorubicin and mAbs alone (Figure 4(a)). Regarding the canine OSA cells, as shown in Figure 4(b), the addition of doxorubicin...
at the lowest selected dose (100 nM) resulted per se effective in significantly inhibiting tumor cell proliferation, however a significantly higher inhibition was observed when the combination of mAbs and doxorubicin was applied, being superior to both single agents alone. As we have previously demonstrated the clinical effectiveness of in vivo CSPG4 immune-targeting by means of DNA vaccination in dogs affected by MM,25,26 to evaluate the potential of this strategy for the treatment of OSA, we incubated human and canine OSA cells with sera collected from canine MM patients enrolled in our previous veterinary trial. The sera were collected at the moment of the surgical removal of the primary tumor (pre-Vax) or after the vaccination (post-Vax) with a plasmid coding for the Hu-CSPG4, that induces the production of anti-CSPG4 antibody in the vaccinated dogs, against both the human and the canine CSPG4 protein.25 Interestingly, post-Vax sera were able to inhibit the proliferation of both human and canine OSA cells as compared with pre-Vax sera (Figure 4(c) and (d)). Together, these results suggest that CSPG4 may have a key role in both human and canine OSA cell proliferation, which could be impaired by its immune-targeting. Of note, our data propose
the potential impact of adjuvant anti-CSPG4 DNA vaccination, alone or in combination with doxorubicin (Supplemental Figure S3) for the treatment of OSA patients, as we previously demonstrated for another aggressive and CSPG4-positive disease, MM.25,26

CSPG4 immune-targeting significantly inhibits CSPG4-positive human and canine osteospheres.

OSA is among the several cancer types in which a relevant role of CSC in tumor initiation and progression is evident. Indeed, CSC are thought to be the main drivers of OSA-related death, being responsible for tumor chemoresistance, finally resulting in recurrence and metastasis.40 Starting from this assumption, we generated MG-63 (Supplemental Figure S4(a)) and Penny (Supplemental Figure S4(b)) derived osteospheres, following the protocol described by Conti and coworkers.33 Interestingly human (Supplemental Figure S4(c)) and canine (Supplemental Figure S4(d)) osteosphere-derived cells expressed high levels of CSPG4, making it an even more interesting antigen for the immune-targeting of both differentiated cancer cells and...
CSC in OSA. Of note, both human (Figure 5(a)) and canine (Figure 5(b)) osteospheres incubated with anti-CSPG4 mAbs pool (225.28, TP32, TP49, and VF20-VT87.41) displayed reduced spheroid viability as compared with control, with each mAb clone showing a similar effect (Supplemental Figure S2(c) and (d)). In addition, we measured the in vitro growth of osteospheres when incubated with doxorubicin alone or in combination with anti-CSPG4 mAbs. For the human MG-63-derived osteospheres we selected the 10 µM doxorubicin dose, since the lower dose was not even effective on epithelial cells (Figure 4(a)). CSPG4 targeting with mAbs pool significantly increased the inhibitory effect of doxorubicin on osteospheres viability (Figure 5(a)). Regarding the canine Penny-derived osteospheres, the 100 nM doxorubicin dose, successfully used for epithelial canine Penny cell proliferation studies (Figure 4(b)) was not effective alone against osteospheres (Figure 5(b)), confirming previous findings that CSC avail of increased chemoresistance ability.41 Using a higher doxorubicin concentration (1 µM) on canine Penny-derived osteospheres, we observed that doxorubicin alone in this case significantly inhibits the sphere viability (Figure 5(b)), however, the combinatorial approach of doxorubicin...
plus anti-CSPG4 mAbs further improved the effect of the single treatments alone (Figure 5(b)). Interestingly, post-Vax sera derived from canine MM patients vaccinated with the Hu-CSPG4 plasmid resulted effective in inhibiting human and canine osteosphere viability, as compared with pre-Vax sera (Figure 5(c) and (d)). Overall, these findings suggest that CSPG4 immune-targeting with a mix of anti-CSPG4 mAbs, and most interestingly, with anti-CSPG4 DNA vaccination, could also have an impact on the CSC compartment, which is considered endowed of a high metastatic potential.

CSPG4 immune-targeting significantly inhibits CSPG4-dependent human and canine OSA cells migration.

Migration is a critical property of cancer cells, which could determine their metastatic behavior. As metastasis is one of the major challenges in OSA treatment, we evaluated the potential of CSPG4 immune-targeting in counteracting the migratory ability of human and canine OSA cells. Interestingly, when cells were pre-incubated with anti-CSPG4 mAbs pool (225.28, TP32, TP49, VF20-VT87.41), a significantly reduced migratory potential, as compared with control, was evident for both human MG-63 (Figure 6(a)) and canine Penny (Figure 6(b)) OSA cells. Moreover, cell motility was also affected by doxorubicin; indeed, a significant migratory reduction was observed when MG-63 (Figure 6(a)) and Penny (Figure 6(b)) cells were treated with 10 µM and 100 nM doxorubicin, respectively. However, the combinatorial approach using doxorubicin plus anti-CSPG4 mAbs pool further improved the effect of the single treatments alone in both human MG-63 (Figure 6(a)) and canine Penny (Figure 6(b)) cells, making the possibility of chemotherapy and adjuvant CSPG4 immune-targeting an even more appealing strategy to fight against the metastatic disease.

Discussions

CSPG4 is a transmembrane protein involved in several protumorigenic signaling pathways. It is expressed in a wide range of highly aggressive tumors, including MM, triple-negative breast carcinomas, leukemia, gliomas, in which it is associated with those hallmarks linked to tumorigenesis including proliferation, invasion, and metastasization. The overexpression of CSPG4 on CSC in different tumor histotypes, could suggest its potential implication also in providing a survival advantage to this subpopulation, considered responsible for recurrences and metastasis. Therefore, for all these properties, the CSPG4 is considered an ideal and safe oncoantigen for anticancer targeted therapies, being
barely expressed on normal healthy tissues. Indeed, by means of an IHC analysis of a FDA Standard Frozen Tissue Array, including 30 different organs, Rivera et al. demonstrated that no CSPG4 expression was found in healthy tissues. In general, a limited CSPG4 expression is associated with stem cells and adult progenitor cells, which however have been suggested to lose its expression during terminal differentiation. Moreover, a heterogeneous CSPG4 expression has been detected on activated pericytes but interestingly, only poorly stabilized vascular structures contain CSPG4 expressing pericytes, whereas CSPG4 is downregulated in pericytes associated with quiescent vessels, and absent or not detectable in pericytes of stable vessels in the adult healthy tissues. On the basis of these considerations, several immunotherapeutic approaches against CSPG4 for the treatment of melanoma and other CSPG4-expressing tumor histotypes have been tested both in preclinical and clinical settings. CSPG4-specific chimeric antigen receptors (CAR) T cells, as well as sophisticated mAb-based approaches have been generated, demonstrating the antitumor potential of CSPG4 immune-targeting. An alternative approach is active immunization, which has demonstrated to bring about effective and long-lasting antitumor responses, without the risk of resistance development. In this direction, some evidence of the effectiveness of active immunization against CSPG4 in melanoma patients was found through vaccination with the anti-idiotypic antibody MK2-23, which bears the internal image of the mAb 763.74 against a defined CSPG4 epitope. Interestingly, the induction of CSPG4-specific antibodies in immunized patients was associated with significantly longer survival and metastasis regression. However, this approach never ended up in clinics, owing to both the difficulties in standardization of MK2-23 and to side effects associated with Bacille Calmette–Guerin administration, the adjuvant required to break immune tolerance and to induce an efficient immune response. However, these encouraging data provided a strong rationale for the development of new strategies of active immunization against CSPG4.

Recently, DNA-based vaccines have raised interest as a concrete and viable anticancer strategy. In this direction, we have recently focused our attention on the antitumor potential of in vivo electroporation of a DNA vaccine (electrovaccination) coding for the Hu-CSPG4 protein. To confer a high translational power to our study, we tested the safety, immunogenicity, and clinical efficacy of the vaccine in prospectively enrolled client-owned dogs with en bloc surgically resected stage II and III CSPG4-positive spontaneous oral MM. The results obtained in our studies demonstrated the ability of the xenogeneic DNA electrovaccination against CSPG4 to break the immune tolerance in dogs and to induce a specific humoral response which relates favorably with a significant prolongation of disease-free and overall survival time in vaccinated dogs with surgically resected MM as compared with controls treated with surgery alone. These results lay the foundation for the evaluation of this immunization strategy for the treatment of other CSPG4-expressing tumors.

To date, OSA still represents a critical challenge in the oncology field, because conventional therapies have demonstrated partial effectiveness only in patients affected by localized tumor, while failing in the treatment of advanced patients. Several strategies have been evaluated to improve the survival of OSA patients without encouraging results. Some clinical trials involving tyrosine kinase targeted therapies or checkpoint inhibitors have been assessed, however, considerable improvements in patients’ outcome have not been realized at all. Therefore, it is clearly evident the urgent need for novel and effective therapies. In this panorama, the identification of CSPG4 as a potential OSA-associated target could offer new possibilities for the treatment of this disease.

For this purpose, in the present study, we first focused our attention on the evaluation of CSPG4 expression in human OSA. We have analyzed mRNA levels for CSPG4 in previously published genome-wide expression data of osteoblasts, MSCs, and 84 high-grade OSA pretreatment biopsies. We detected the overexpression of CSPG4 mRNA in human high-grade OSA biopsies, as compared with the hypothesized OSA progenitors, and validated the CSPG4 protein overexpression in human OSA cell lines. We availed of the collected information regarding the clinical evolution of the disease of the 84 high-grade OSA patients included in the dataset. All patients underwent neoadjuvant chemotherapy, and for all patients, the Huvos necrosis grading system was applied for the assessment of chemotherapy efficacy. Unfortunately, patients with a poor histologic response to neoadjuvant chemotherapy (mostly Huvos Grade I–II) showed no
benefit following distinct postoperative therapies. \[^{49}\] Interestingly, CSPG4 mRNA was found overexpressed independently from the Huvos Grade and a higher degree of expression resulted in lower grades. We further evaluated whether the CSPG4 expression is related to OSA patients’ prognosis. Actually, OSA patients show a shorter overall survival and metastasis-free survival probability when CSPG4 is overexpressed. Therefore, taken as a whole, these results suggest the strong potentiality of adjuvant CSPG4 targeting for both good and poor responders to chemotherapy.

In addition, because of the age of its onset, OSA is socially important and limits the possibility of testing new therapies; the identification of reliable models of human OSA is therefore a critical point. For this reason, we decided to evaluate the feasibility of using spontaneous canine OSA as a preclinical model to test anti-CSPG4 immunotherapies. Indeed, canine patients spontaneously develop tumors as humans do, in a context of an intact immune system, with strong anatomical and physiological similarities with the human counterpart. This is true also for OSA. Moreover, as in humans, treatments for canine OSA includes mainly surgery and chemotherapy, which however are often disappointing with only 20% of canine OSA patients treated using the current standard of care still alive at 1 year and lung metastasis being the most relevant cause of death. \[^{50-52}\] A tangible example of the importance of comparative oncology in OSA came from the early 1990s, when limb-sparing methods pioneered in canine patients with OSA \[^{83}\] have become standard of care for human patients, which today clearly benefit from advances made in both surgical treatment and in the provision of supportive care.

With this in mind, we have demonstrated that CSPG4 expression is detectable, with variable expression levels, in a high percentage of canine OSA biopsies by means of IHC. In addition, also canine OSA cell line showed to highly express CSPG4, representing an interesting tool to be exploited for in vitro studies. As for humans, the Kaplan–Meyer curves suggest that CSPG4 overexpression is also related to a poor prognosis in canine patients. Thus, with this study we highlighted the potential clinical relevance of evaluating anti-CSPG4 strategies in canine OSA patients, with a high translational value. For this reason, we examined the potentiality of CSPG4 immune-targeting against both human and canine OSA cell lines in vitro, in order to consider anti-CSPG4 immunotherapy as a potential new weapon against OSA.

CSPG4 is implicated in several cellular processes, \[^{24,39}\] therefore its targeting could impair simultaneously different steps in the tumorigenic process. First, our results suggest that CSPG4 is involved in OSA cell proliferation. Indeed, we showed that four anti-CSPG4 selected mAbs (225.28, TP32, TP49, and VF20-VT87.41) are able to significantly impair the proliferation of both human and canine OSA tumor cells. This inhibition is evident when mAbs are used in a mixed pool or when used as single agents, suggesting that the engagement of different antigen epitopes by each clone does not seem to have different effects on cancer cell survival. The effect of CSPG4 immune-targeting in vitro is evident although modest. This can be attributed to several reasons: both OSA cell lines are not 100% positive for CSPG4 expression, therefore there will be a CSPG4-negative population able to escape to mAbs treatment; the selected dose of mAbs used is low, so better effects could be achieved increasing the dose. However, an interesting finding is the ability of mAbs treatment to significantly sensitize OSA cells to doxorubicin. Actually, the combination of doxorubicin, one of the most common chemotherapeutic agent used in both human and veterinary setting, \[^{41}\] with anti-CSPG4 mAbs enhanced the inhibition of cancer cells’ growth. Of note, on one side CSPG4 can regulate the AKT–pAKT pathway considered responsible for chemoresistance, on the other side we have previously demonstrated that anti-CSPG4 mAbs can induce a downregulation of the CSPG4 receptor when incubated with CSPG4-positive melanoma cells, \[^{26}\] and this could consequently impair the downstream signaling, leading to the reduction of the AKT–pAKT axis (unpublished data and Rolih et al. \[^{24}\]). Therefore, the association of chemotherapy and anti-CSPG4 immune-targeting open up the possibility of increasing the anti-tumor effect of single agents alone, combining standard of care with novel strategies.

Moreover, the intrinsic OSA chemoresistance may be the result also of a privileged survival of a population of tumor cells, that is, CSC to which are associated tumor recurrence and metastasis development following chemotherapy. For this reason, in this study we evaluated the ability of mAbs alone or in combination with doxorubicin to impair not only epithelial cancer cells but also osteospheres enriched in CSC. First, we demonstrated that
subsequent passage of CSC-enriched osteospheres retain the CSPG4 overexpression, making it an interesting antigen to target CSC too. Then, we showed the ability of mAbs, alone or in combination with doxorubicin, to inhibit osteospheres viability. Also in this case, the selected anti-CSPG4 mAbs (225.28, TP32, TP49, VF20-VT87.41) showed a similar effect when used together in a pool or when tested individually. These results propose the potentiality of CSPG4 immune-targeting for the elimination of CSC and the prevention of recurrences and metastasis in OSA. Moreover, our data suggest a potential involvement of CSPG4 in OSA cell migration, because we demonstrated the significant impact of CSPG4 immune-targeting against the migratory ability of both human and canine OSA cell lines, highlighting the pleiotropic effects of anti CSPG4 mAbs. In addition, mAbs treatment significantly increases the antimitagatory effect of doxorubicin, as demonstrated by Transwell assays, supporting the relevant clinical consequence of combinatorial anti-CSPG4 immune-targeting and chemotherapy to fight against OSA metastasis. Overall, these results provide an additional step forward in the understanding the impact of CSPG4 in its whole for OSA progression.

Finally, to consider anti-CSPG4 DNA electrovaccination as a new therapy for the adjuvant treatment of OSA, on the basis of our previous positive results obtained for MM, we used sera derived from MM canine patients adjuvantly treated with Hu-CSPG4 DNA plasmid to evaluate the ability of vaccine induced antibodies to inhibit the proliferation of OSA cells and osteospheres in vitro. Interestingly post-vaccination sera were effective in inhibiting cell growth and sphere viability alone or in combination with doxorubicin. These results suggest the potential efficacy of our DNA vaccination strategy also for the treatment of canine OSA patients in vivo, with a strong translational value for human OSA management.

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Author contributions
FC, PB, and FR contributed to the conception and design of the study. FC, FR, and LT wrote the manuscript. FR and LT performed the in vitro experiments. SI performed the immunohistochemical analyses. DG, FS, EM, MM, FG, and PB contributed to patient recruitment and clinical follow up. MA performed the meta-analysis and data collection from human datasets. RDM and SF provided reagents and cell lines. All authors contributed to collection, analyses and interpretation of data, reviewed, and approved the final submitted version.

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Conflict of interest statement
The authors declare that there is no conflict of interests.

Supplemental material
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References


