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Cellular and Phenotypic Characterization of Canine Osteosarcoma Cell Lines

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Abstract

Canine and human osteosarcoma (OSA) have many similarities, with the majority of reported cases occurring in the appendicular skeleton, gender predominance noted, high rate of metastasis at the time of presentation, and a lack of known etiology for this devastating disease. Due to poor understanding of the molecular mechanisms underlying OSA, we have characterized seven different OSA canine cell lines: Abrams, D17, Grey, Hughes, Ingles, Jarques, and Marisco and compared them to U2, a human OSA cell line, for the following parameters: morphology, growth, contact inhibition, migrational tendencies, alkaline phosphatase staining, heterologous tumor growth, double-strand DNA breaks, and oxidative damage. All results demonstrated the positive characteristics of the Abrams cell line for use in future studies of OSA. Of particular interest, the robust growth of a subcutaneous tumor and rapid pulmonary metastasis of the Abrams cell line in an immunocompromised mouse shows incredible potential for the future use of Abrams as a canine OSA model. Further investigations utilizing a canine cell model of OSA, such as Abrams, will be invaluable to understanding the molecular events underlying OSA, pharmaceutical inhibition of metastasis, and eventual prevention of this devastating disease.

Key words: Abrams, Canine, Osteosarcoma, U2

Introduction

Osteosarcoma (OSA) is a deadly form of musculoskeletal cancer which develops from the cells responsible for forming the bone matrix.²² In children and adolescents, OSA is the most common type of primary bone cancer,^{5,23} with approximately 1-3 cases per million annually, and 70-75% of patients being 10 to 25 years of age.^{12,19} Interestingly, the incidence of OSA is slightly higher in males than females, with a ratio of 1.5:1.^{19,23} Factors underlying the development of OSA are still largely unknown, however humans are at a higher risk following periods of rapid adolescent growth, with the metaphyseal area adjacent to the growth plate of long bones most commonly affected.^{19,23} Most OSA start as solitary lesions, especially within the distal femur, proximal tibia, and proximal humerus.²² Although micrometastatic disease is likely present in approximately 80% of human patients, it is only detected in 8-15%,^{7,8} and as a result patients often die from pulmonary metastatic disease.^{5,22}

In canines, OSA is more common than it is in humans, with approximately 10,000 new cases each year.³ OSA is generally a disease of large breed canines where increasing height is a more prominent risk factor then body weight.¹⁸ As in humans, males more commonly present with OSA than females,^{5,10} although some studies reveal no prevalence between sexes.¹ When a dog presents with OSA, less then 10% of the cases have radiographic evidence of lung metastasis.²² Canine OSA is most often found in the appendicular skeleton,^{5,8,24} originates in the metaphyseal region of the long bones (similar to humans), and occurs twice as frequently in the thoracic versus the pelvic limbs. The most common site of canine OSA is the distal radius followed by the proximal humerus.^{5,18} Despite treatment with amputation and chemotherapy, over 80% of dogs succumb to metastatic disease within 2 years of diagnosis.^{5,24}

To summarize, many commonalities between human and canine OSA exist: the long bones are most often affected, males are affected more commonly than females, the etiology is unknown, and very few cases show lung metastasis at the time of presentation. Because of these similarities, the characterization and use of a canine model of OSA could be invaluable in understanding this devastating disease, aiding in assessing potential therapeutic agents and potentially preventing disease formation. In this report, we have characterized six canine OSA cell lines for morphology, growth, contact inhibition, migrational tendencies, alkaline phosphatase protein, heterologous tumor growth, double-strand DNA breaks, and oxidative damage and compared them to U2, a human OSA cell line. Further investigations utilizing a canine cell model of OSA with xenograft potential will be invaluable to understanding the cellular and molecular events underlying OSA.

Materials and Methods

Cell Culture: Canine OSA cell lines Abrams, D17, Grey, Hughes, Ingles, Jarques, and Marisco were a generous gift from Dr. Greg MacEwan, University of Wisconsin, Madison, Wisconsin. The human cell line U2, was purchased from ATCC. All OSA cell lines were grown in Modified Eagles Medium (MEM; CellgroTM) supplemented with 15% fetal bovine serum (FBS; HyCloneTM), 2% MEM vitamins, and 1% each L-glutamine, non-essential amino acids, and so dium pyruvate. All cells were maintained at 37° C, humidified with 5% CO₂.

Cell Proliferation: Abrams, D17, Grey, Hughes, Ingles, Jarques, and U2 were used in this experiment. In order to determine the varying proliferation rates of the cell line(s), a seven-day proliferation trial was performed. To take into account differences in cell size, as some OSA cells were quite large and spindloid in comparison to other cell lines, cells were harvested at the onset of this experiment when flasks reached 90% confluency and then each cell line was diluted similarly before being added to the 100 mm² plates. Thus, each 75 cm² flask was trypsinized when cells were 90% confluent with 1 ml 0.25% trypsin-EDTA 0.25% for 5 min. MEM with 15% FBS (22 mL total volume) was added to each flask and the cells were thoroughly mixed. One mL of this suspension was removed and cells were counted to establish the count at day 0. One mL aliquots of the remaining 21 mL were added to 9 mL of MEM with 15% FBS in 100 mm² plates. Cell counts were performed daily (22 - 24 h intervals) as follows: 3 plates from each cell line were trypsinized with 0.5 mL 0.25% trypsin-EDTA for 5 min, 3 mL MEM with 15% FBS was added, and all 3 dishes were combined and thoroughly mixed. Each sample was diluted 1:10 with 0.4% PBS and 37% formaldehyde solution and counted twice. This resulted in the number of cells/mL using the following calculation: [(value from counter #1) + (value from counter #2] X 10 = cells/mL. In order to standardize the cell count data the cell number of subsequent days was divided by the number of cells counted on day zero. Counting was performed using a Particle Data Counter (El Zone 180 Particle Data, Inc) and double-checked using a hemocytometer. The experiment was performed in triplicate for each cell line for 7 d.

Growth in Soft Agar: Abrams, D17, Grey, Hughes, Ingles, and U2 were used for this experiment based on their performance in the cell proliferation experiment above. Agar (GibcoTM) was reconstituted with 100 mL MEM with 15% FBS to 0.3% and 0.8% final concentration, boiled for 1 min and autoclaved. The agar was cooled slightly, then 100 mL of agar was mixed with 2.5×10^5 cells in 5 mL MEM with 15% FBS. Each cell-agar mixture (5 mL) was added to a 100 mm² dish. Each experiment was performed in duplicate. The plates were visualized at 4x daily and colony measurements were taken and photographed on 4X power at 8d, 10d, 11d, 15d, 18d and 22d post-plating. Images were acquired using a Zeiss Axiovert 200M microscope equipped with a 63X 1.4 N/A oil immersion objective and Hammatsu ORCA-ER cooled charge-coupled device camera.

Boyden Chamber Assay: Abrams and U2 were assessed in this experiment due to the strong performance of Abrams in the cell proliferation and colony growth experiments. Ingles was also used as a poor growth control cell line (data not shown). The cell lines were grown to 80% confluency and FBS starved for 24 h prior to harvesting. Cultures were harvested and volume adjusted to yield 2.5×10^6 cells/mL. Cells suspended in either serum-free growth medium (300 µL) or normal growth medium (300 µL) with were added to the appropriate wells (fibronectin- or BSA-coated) of the Boyden chambers under sterile conditions, as per manufacturers instructions. The

cells were incubated for 24 h and 6 drops of Cell Stain Solution were added to each well after removal of The fibronectin-coated Boyden chambers media. from the test plate were carefully removed and the cells were removed from inside the chamber. Boyden chambers were then placed in the test plate wells with stain (from above). The same procedure was followed for the BSA-coated control chambers and they were placed in the control wells with stain. Plates were then incubated at room temp for 30 min. Cell stain was removed and the wells were washed with PBS. Extraction buffer (300 µL) was added to the empty wells in the test and control plates. Boyden chambers were washed thoroughly with distilled water, cleaned and placed into extraction buffer in their respective plates. Plates were gently shaken to elute stain for 10 min, and then 100 µL of stained solution was transferred from the elution wells into a microtiter plate. Absorbance at 540 nm was analyzed.

Heterologous Tumor Growth and Metastasis: Abrams, Ingles, and U2 were used for this experiment. Rag1 male mice were purchased from the Jackson Laboratory, and acclimated in Colorado State University's central animal care facility, which is fully accredited by the American Association for Accreditation for Laboratory Animal Care. All animal procedures were conducted following an Animal Care and Use Research Protocol approved by the Colorado State University Animal Care and Use Committee. Mice were allowed ad libitum access to food and water throughout the duration of experimentation. 8 wk old mice were injected subcutaneously mid-dorsally with 1 x 10⁶ cells suspended in 0.2 mL PBS, and monitored daily for tumor growth. At 10 wk post-injection, a MRI was performed on each mouse to determine possible lung metastasis. Mice were then euthanized, necropsied, and tissues submitted for histopathologic examination.

Alkaline Phosphatase Staining: Abrams, D17, Grey, Hughes, Ingles, Jarques, and U2 were evaluated for expression of alkaline phosphatase and fibroblasts were used as negative controls. Chamber slides were fixed for 10 min in 1% paraformaldehyde and air-dried. Slides were rehydrated in buffer (100mM Tris-HCl, 100mM NaCl and 50mM MgCl₂), washed in TBS with 0.05% TBS (TBS-T0.5) and coated with DI water to keep moist. For staining, 100µl BCIP/NBT, prepared as instructed in kit (Vector Laboratories, SK-5400) was added to each reservoir without introducing bubbles and slides were allowed to sit covered and in the dark for 1 h. The slides were rinsed with TBS-T0.5 and another 100 µl BCIP/NBT was added to each reservoir. The slides were incubated overnight and 100 µl BCIP/NBT was added to each reservoir, allowed to sit for 1 h and rinsed with TBS-T0.5. The slides were then counterstained with hematoxylin for 30 s and coverslipped with Kaiser's aqueous mounting media.

Phosphorylated H2AX Immunocytochemistry: Cells were cultured for 24 h on plastic chamber slides and then washed with PBS followed by fixation with 4% paraformaldehyde for 15 minutes. Following a second wash with PBS, cells were then permeabilized with 0.2% Triton X with PBS for 10 minutes. Cells were then covered with parafilm and incubated with 10% goat serum with PBS overnight at 4°C. Following overnight incubation, the cells were then covered with parafilm and incubated with a mouse monoclonal phosphorylated histone H2AX (Ser139) (Millipore) in 10% goat serum with PBS for 1 hour at 37°C. The cells were then washed three times for 10 minutes each in PBS, followed by a 1-hour incubation at 37°C with the appropriate secondary antibody. The slides were then washed three times for 10 minutes each in PBS, and then counter stained with Vectashield mounting media with DAPI. The cells were cover slipped and visualized with Olympus AX70 equipped with a cooled CCD Sensys camera. MacProbe was utilized to obtain images.

Oxidative Damage: Cells were cultured for 24 h on plastic chamber slides and then washed with PBS followed by fixation with 4% paraformaldehyde for 15 minutes. Following a second wash with PBS, cells were then permeabilized with 0.2% Triton X with PBS for 10 minutes. Cells were then covered with parafilm and incubated with 10% goat serum with PBS overnight at 4°C. Following overnight incubation, the cells were then covered with parafilm and incubated with a FITC-8-oxoguanine protein conjugate (Kamiya Biomedical Company) in 10% goat serum with PBS for 1 hour at 37°C. The cells were then washed three times for 10 minutes each in PBS and counter stained with Vectashield mounting media with DAPI. The cell is cover slipped and visualized with Olympus AX70 equipped with a cooled CCD Sensys camera. Mac-Probe was utilized to obtain images.

Statistical Methods: Analysis of variance was used to determine statistical significance with GraphPad Prism software. When statistical differences were detected, Tukey's *post hoc* test was used to determine individual variations. As the U2 cell line is currently employed most commonly and the purpose of these experiments is identifying whether a comparable canine cell line exists, all means were compared to U2.

Results

During the 7-day testing period, the cell lines

were vastly different in their proliferation characteristics (Fig. 1). Overall cellular proliferation demonstrates that at day 7 the Abrams line was an aggressively growing OSA cell line, as it neither died off nor reached a plateau. Additionally, the Abrams line quickly grew colonies in soft agar, an indicator of anchorage-independent growth, and produced one of the highest colony diameters, 2768.9 µm over the 22 day testing period (data not shown). In comparison, maximal diameters for the other cells lines were as follows: 2339.3 µm for D17, 1301.6 µm for U2, and 529.3 µm for Ingles. Ingles in addition to having minimal soft agar colony growth over the 22 day period was also one of the slower proliferators, to the point where the data needed to be represented on a different Y-axis in order to be visualized (Fig 1). The cell lines Grey and Hughes were suicidal early in the cell proliferation assay, dying by day 4 in the assay (data not shown), and had the smallest colony growth in soft agar after 22 days (529 µm maximal diameter for both of these lines). Additionally, Jargues also grew poorly in the proliferation assay (data not

shown as it was lower than Ingles), was difficult to maintain in culture and did not grow in the soft agar. Thus, as we were looking for a cell line that would model canine osteosarcoma over usage of the human U2 cell line, Grey, Hughes and Jarques were not carried further in the Boyden Chamber assay nor the xenografting experiment.

The migrational capacity of cells is indicative of metastatic potential. Boyden chamber wells were coated with FN to assess migration or BSA as a negative control. Abrams demonstrated significant migrational tendencies compared to the BSA negative controls (Fig. 2) and performed significantly better than U2 in this assay. Ingles, as the negative or poor growth control demonstrated no migrational tendencies (data not shown). Furthermore, the Abrams cells consistently stained positive for alkaline phosphatase (Fig. 3-4) with all other cell lines having a poor and inconsistent staining pattern. Based upon these results, Abrams was selected as an aggressive OSA line comparable to U2.



Figure I. OSA cell lines proliferate differently over time, with the Abrams cell line outperforming U2. Cell numbers per cell line (U2, Abrams, D17 and Ingles represented) over the 7 day experimental period. Please note that the cell numbers (on the Y axis) spans 0-200,000 cells for all cell lines except the Ingles which proliferated so slowly that the Y-axis spans 0-1,000 in order to visualize the data for this cell line. Error bars represent standard deviation.



Figure 2. Abrams cells migrate more extensively than U2 cells in the Boyden chamber. Chamber wells were coated with fironectin (Black Bars) or bovine serum albumin (Checkered Bars), and 750,000 cells were added to each well. After 24h cells were stained, lysed, and optical density at 540 nm was read. Statistical differences were determined via two-way analysis of variance, with cell type (p=0.003) and coating (p<0.001) being significant, as well as the cell type and coating interaction (p=0.0086).

Subcutaneous injection of Abrams cells resulted in tumor growth in the skin with lung metastases 10 wks post-injection (Fig. 3). These tumors were grossly evident in the skin and could be identified grossly as suspicious areas within the lung tissue and histopathologically in both the skin and lung. However, use of MRI at 10 weeks was not sensitive enough to declare metastasis and only after going back to the images after gross and histological examination could the areas be characterized as suspicious on MRI scan. Other internal organs were examined grossly and histopathologically and did not contain evidence of metastases. Subcutaneous injections of Ingles or U2 cells did not result in tumor growth in the skin nor metastasis to lungs or other organs.

 γ H2AX immunocytochemistry demonstrated that the biologically more aggressive canine and human OSA cell lines (Abrams, D17, Grey, Hughes, Marisco, and U2) demonstrated more nuclear foci of phosphorylated γ H2AX as compared to the less aggressive canine OSA cell lines (Ingles and Jarques) (Fig. 4).

Additionally, the biologically more aggressive canine OSA cell lines (Abrams, D17, Grey, and Marisco) demonstrate increased oxidative damage as evidenced by increased mitochondrial and nuclear staining of 8-oxoguanine. The canine OSA cell line Hughes demonstrates intermediate staining, while the less aggressive canine OSA cell lines (Ingles and Jarques) and the human OSA cell line (U2) demonstrate minimal staining (Fig. 5).



Figure 3. The Abrams **OSA** cell line produces an aggressive tumor in the Ragl mouse. I. Heterologous tumor growth of Abrams cells in Rag1 mouse. Grossly, the tumor measures 13.4 mm long x 8.8 mm wide 10 wk post injection. 2. Histopathology of heterologous Abrams skin tumor 10 wk post xenografting via subcutaneous injection (Hematoxylin and Eosin). Arrow denotes a mitotic figure. 3. Heterologous Abrams cell tumor metastasis to lung (Hematoxylin and Eosin). Inset of lung metastasis at higher power. 4. Abrams cells stain positively for alkaline phosphatase with hematoxylin counterstain.







Figure 5. The biologically more-aggressive OSA cell lines demonstrate increased mitochondrial and nuclear oxidative evidence characterized by increased DNA 8-oxoguanine. In all cell types mitochondrial and nuclear 8-oxoguanine foci are stained green with DAPI for nuclear counterstain. Abrams, D17, Grey, and Marisco demonstrate increased oxidative damage as evidenced by increased mitochondrial and nuclear staining of 8-oxoguanine. Hughes demonstrates intermediate staining, while the less aggressive canine OSA cell lines (Ingles and Jarques) and the human OSA cell line (U2) demonstrate minimal staining of 8-oxoguanine. Representative images from the following osteosarcoma cell lines are within the panels indicated: A) Abrams, B) D17, C) Grey, D) Hughes, E) Ingles, F) Jarque, G) Marisco, H) U2.

Discussion

Although many common characteristics exist between human and canine OSA, the underlying etiology of the disease is poorly understood in either species. Initial genetic characterization of a few human and canine OSA cell lines have indicated mutations in PTEN16, inactivation of p5315 and overexpression of met and sis oncogenes6,17 to be of importance in these tumors. Further characterization of canine OSA cell lines will provide information with which to select an appropriate in vitro model to aid in identifying molecular and/or genetic mechanisms underlying this devastating tumor, as well as provide an experimental platform to test potential therapeutic interventions. Based upon our results, the canine osteosarcoma Abrams cell line could indeed be utilized as such, as these cells possess more characteristics indicative of metastatic OSA compared to the human cell line currently employed, U2.

Of all lines tested, Abrams continued its rapid proliferation without dying off, indicative of an aggressive and continuously growing phenotype. Further, the Abrams cells showed significant differences in migration potential. Additionally, the Abrams cell line produced large colonies in soft agar, indicating capability of anchorage-independent growth with little contact inhibition, indicative of carcinogenesis and metastatic potential. Only the Abrams cells consistently stained positively for alkaline phosphatase, an enzyme used to differentiate OSA from other soft tissue tumors on histopathology, and one that is often used to correlate with prognosis in OSA patients.7,11,20,21 In some studies, elevated serum levels of alkaline phosphatase are correlated with approximately 50% shorter survival times despite assertive treatment.7,11,19

Previous experiments correlate significant cell migration in a Boyden chamber with increased invasive properties in vivo.4 The Abrams cells migrated extensively, demonstrating a high potential to be invasive in vivo, surpassing even the migrational capacity of the human U2 line. Following subcutaneous injection into Rag1 mice, only the Abrams cell line formed subcutaneous tumors at the site of injection and metastasized to lung tissue. Fitzpatrick et al. demonstrated that the D17 cell line could be utilized as a xenograft, but there was no indication of metastasis (9). Moreover, Kanaya et al. utilized the POS cell line in their studies of the anti-tumor effect of adenoviral-vector mediated p53 gene transfers in nude mice but there again was no indication of metastasis after POS xenograft formation in these mice.¹³ As tumor development and subsequent formation of metastases

are hallmark characteristics of OSA, the Abrams cell line proved to be a good choice for a model of OSA metastasis. Osteoid was not easily detectable in histopathologic sections of the skin or lung tumor sites, though it is considered to be a histological indicator of OSA. However, minimal or non-detectable osteoid may be secreted in poorly to undifferentiated OSA and thus not be identified on routine histological section. Moreover, additional characteristics of neoplasia were noted including open nuclei, frequent mitotic figures, anisokaryosis, and anisocytosis (Fig. 3).

Carcinogenesis in general is associated with DNA replication stress and oxidative damage, resulting in DNA double-strand breaks, genomic instability, and selective pressure for p53 mutations. DNA double strand breaks result in massive relaxation of chromatin coiling.14 Misrepaired or unrepaired DNA double strand breaks contributes to chromosomal aberrations leading to cell death, mutation, and oncogenic transformation.14 H2AX is a member of the histone H2A family and is involved in nucleosomal organization of chromatin.² The basic subunit of chromatin, the nucleosome, consists of a core of eight proteins, two from each of the H2A, H2B, H3, and H4 families, with about 140 bp of DNA coiled around the core and the fifth histone family, H1, on the linker DNA acting as a bridge between adjacent nucleosomes.² Each nucleosome contains two H2A molecules and H2AX is synthesized in a replication-dependent and replication-independent manner.² Upon double-strand break formation, members of the PI3K-like kinase family, including ataxia telangiectasia mutated (ATM), ataxia telangiectasia and Rad3-related (ATR), and DNA-dependent protein kinase (DNA-PK), are activated and phosphorylate H2AX and other DNA repair and checkpoint proteins.² As double-strand DNA breaks are repaired, yH2AX foci disappear.² There are two potential mechanisms of yH2AX removal: dephosphorylation of yH2AX or removal of phosphorylated yH2AX from the chromatin.² Dephosphorylation depends upon the phosphatases PP2A and PP4C. Phosphorylation of the H2AX histone variant on a serine four residues from the carboxyl terminus is a very sensitive assay for double strand break measurements.^{2,14} Thirty minutes following a DNA double strand break, large numbers of yH2AX molecules form in the chromatin around the break site, creating a focus where proteins involved in DNA and chromatin remodeling accumulate.² The Abrams cell lines demonstrate high levels of both double-strand DNA breaks and oxidative damage, further supporting its use as an aggressive canine OSA cell line.

The growth of cancers is accompanied by progressive infiltration, invasion, and destruction of the surrounding tissues. Given how quickly the Abrams cell line proliferated, its ability to migrate in the Boyden chamber experiment, positive staining for alkaline phosphatase, increased numbers of double-strand DNA breaks, increased mitochondrial and nuclear oxidative damage, and lack of contact inhibition in soft agar, the Abrams line is an excellent OSA canine model, particularly metastatic OSA. Because this cancer is so devastating in the canine, further experimentations to develop therapeutic interventions would benefit veterinary medicine. Furthermore, with the myriad of similarities between species, it would be beneficial to have a working canine OSA model that could be used to further the advancement of a human cure. A comprehensive protein comparison between canine and human OSA would allow commonalities to be known and studies undertaken to alter the expression of specific proteins with the goal of tumor suppression. Evaluation of double-strand DNA breaks in spontaneously occurring canine OSA cases may provide additional clinically relevant prognostic information. Given how rapidly OSA metastasizes, and that metastatic spread greatly reduces the possibility of remission of the disease, no achievement would confer greater benefit on patients then methods to prevent the disease and/or prevent its distant spread. Currently, the inhibition of OSA metastasis, as well as the suppression of tumor growth, is still one of the most urgent issues in orthopedic oncology. Identification of canine transplantable OSA cell lines such as Abrams, are potentially useful to clarify factors that are involved in cell-to-matrix interactions as well as for analysis of the aggressive behavior and metastatic potential.

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Conflict of Interest

The authors have declared that no conflict of interest exists.

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