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Histology and Histopathology

From Cell Biology to Tissue Engineering

Ezrin and moesin expression in canine and feline osteosarcoma

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Summary. Biological features of canine osteosarcomas (OS) differ markedly from those found in feline and resemble more human osteosarcomas, in particular for their high rate of metastasis and poor prognosis. Ezrin, radixin and moesin are members of the ERM protein family and link the actin cytoskeleton with the cell membrane. Ezrin and moesin have been shown to be of prognostic significance in tumor progression due to their role in the metastatic process. The objective of this study was to analyze ezrin and moesin protein expression in a series of dog (n=16) and cat (n=8) osteosarcoma samples using immunohistochemistry and western blot techniques. We found that cat OS have a higher moesin expression compared to dog OS, however, the active phosphorylated forms of moesin and ezrin Tyr353 were more abundant in the dog samples. A statistically significant difference was found for the low and high immunohistochemical scores of ezrin and pan-phospho-ERM proteins between cat and dog. Although phosphoezrin Thr567 was higher in feline OS, the membranous localization in dog OS samples indicates the presence of the biologically active form. Therefore, the observed differences in phosphorylated forms of ezrin and moesin status should be further studied to demonstrate if they are relevant for different biological behavior between dog and cat OS.

Key words: Cat, Dog, Osteosarcoma, Ezrin, Moesin, Western blot

Introduction

Osteosarcomas (OS) are the most frequent malignant bone tumors in dogs and cats (Priester and Mc Kay, 1980; Quigley and Leedale, 1983). Treatment options for this neoplasia include surgery, radiation therapy and chemotherapy. However, extended therapies are more usual for dogs than cats, the latter often demonstrating long term control after amputation alone, without any adjuvant treatment (Bitetto et al., 1987). It has been observed that only a minority of cats develop distant metastasis, whereas in contrast the majority of dogs die as a result of metastatic disease (Brodey and Riser, 1969; Dimopoulou et al., 2008). Similar to dogs the osteosarcoma of children and young adults is also highly aggressive and the prognosis is poor in patients with metastasis (Ta et al., 2009). Thus, identifying proteins which are primarily responsible for the development of metastases is particularly important for this disease.

Ezrin, a member of the ERM-(ezrin, radixin, moesin) family, which links the plasma membrane to the cytoskeleton, has been identified as a key component in tumor progression in several tumor types such as breast cancer, gastric cancer or osteosarcoma in humans (Khanna et al., 2004; Jin et al., 2012; Gschwantler-Kaulich et al., 2013). Experimental ezrin overexpression correlated with metastatic progression in human breast cancer cells, whereas ezrin silencing reversed this

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behavior (Li et al., 2008). A significant association between expression pattern and prognosis has been reported, therefore, ezrin is a predictive marker for human osteosarcoma (Kim et al., 2007; Salas et al., 2007; Ferrari et al., 2008; Boldrini et al., 2010). Moreover, ezrin mRNA expression was significantly higher in human osteosarcomas with metastases compared to cases without metastases (Ogino et al., 2007).

OS are classified into several subtypes (osteoblastic, chondroblastic, teleangiectatic, fibroblastic, giant cell rich, multipatterned, poorly differentiated) according to the WHO classification system (Slayter et al., 1994). Ezrin expression has been related to the histological tumor subtype in human osteosarcoma: 30% of osteoblastic osteosarcomas did not express ezrin whereas all other investigated subtypes (fibroblastic, chondroblastic, telangiectatic) were almost generally positive (Ferrari et al., 2008). The relevance of tumor grading and the classification of OS subtypes have also been shown for the canine osteosarcoma (Loukopoulos and Robinson, 2017). To our best knowledge data about abundancy of ERM proteins and correlations with OS subtypes are not available for feline or canine osteosarcoma. A significant decrease of disease-free interval in the case of high ezrin expression has been reported (Khanna et al., 2004). The two functional stages of ezrin that can be discerned within normal and tumor cells are an inactive stage with the N- and C-terminal domains of the protein linked together forming a closed ring-like structure distributed within the cytoplasm, and an opened, active stage where it binds F-actin to the membrane via transmembrane proteins (Tsukita and Yonemura, 1997).

The localization of ezrin should be determined in tumor cells as it is associated with different functional stages of this protein. There was a significant reduction in disease-free survival time probability when ezrin was associated with the membrane in addition to being found in the cytoplasm (Krieg and Hunter, 1992). At least two events have been reported to induce the active ezrin stage - phosphorylation of a conserved threonine (Thr) residue at the aa position 567 and binding of a phosphatidyl inositol-4,5-bisphosphate molecule in socalled FERM domain of the protein (Matsui et al., 1998, 1999; Hamada et al., 2000). Ezrin and its phosphorylated form Thr567 have been demonstrated in human osteosarcoma; however, its permanent expression or overexpression per se was not sufficient for formation of primary tumors or lung metastases. Moreover, a dynamic regulation of ezrin phosphorylation has been suggested for tumor progression and metastasis formation (Ren et al., 2012). At least two other phosphorylation sites have been detected on the ezrin protein associated with its activation: Tyr353 and Tyr477 (Krieg and Hunter, 1992; Bretscher, 1999; Naba et al., 2008). In contrast to phospho-ezrin Thr567, which switches to the plasma membrane upon activation, ezrin phosphorylated at Tyr353 was found to be translocated into the nucleus in osteosarcomas (Di Cristofano et al., 2010). The expression of moesin, another member of the ERM family, has been correlated to cancer progression as well, but has been studied to a much lesser extent compared to ezrin. Increased moesin expression promotes epithelialmesenchymal transition (EMT) in mammary epithelial cells by regulating adhesion and contractile elements for changes in actin filament organization (Haynes et al., 2011). High expression levels were detected in human metaplastic breast carcinomas as well as in nonmammary carcinosarcomas (Wang et al., 2012). Estecha and co-workers have shown that moesin is necessary for lung colonization by melanoma cells, a predominant site for osteosarcoma metastasis as well (Estecha et al., 2009). Even though the etiology of osteosarcoma is not fully known yet, dysregulation of multiple intracellular signaling pathways, especially the phosphatidylinositol 3-kinase (PI3K)/Akt pathway is important for OS onset and progression (for review see Zhang et al., 2014). A large body of evidence suggests that deregulation of this pathway has been involved in multiple pathological processes, including tumorigenesis, proliferation, invasion and metastasis - that means in processes where ERM-proteins involvement has been described as well (Levine et al., 2002; Krishnan et al., 2006; Zhang et al., 2014; Zhu et al., 2014).

There is a striking difference in the frequency of metastasis formation between canine and feline osteosarcoma, although the histopathological patterns are comparable (Dimopoulou et al., 2008). As it has been demonstrated by several investigators that ERM proteins could play an essential role in tumor progression, we hypothesized that there is a difference in ezrin and moesin expression and/or phosphorylation between the highly metastatic canine osteosarcoma and the low metastatic feline osteosarcoma. Moreover, the subcellular localization might vary in these species due to different behavior of the tumor cells. We tested this hypothesis by demonstrating the localization and distribution of ezrin and activated (phosphorylated) stages of ezrin, moesin, Akt and phosphorylated Akt in feline and canine osteosarcoma cells by means of immunohistochemistry and semi-quantitative protein analysis by western blotting.

Material and methods

Tissue samples

Clinical canine (n=16) and feline (n=8) OS samples were collected according to the rules of the ethical committee at the Veterinary University Vienna during therapeutical intervention (amputation or biopsy) or necropsy (animal data are summarized in Table 1). All investigated samples were primary osteosarcomas except one lung metastasis (dog 14) where we did not have access to the primary tumor. The information about the presence of metastases at time of diagnosis is summarized in Table 1. All samples were archived as

formaldehyde-fixed (4%) paraffin embedded (FFPE) specimens and fresh-frozen aliquots were stored at 170°C in the gas phase over liquid nitrogen in the VetBioBank of the Veterinary University Vienna. Samples were decalcified with 8% EDTA before embedding if necessary. Histologic diagnosis and subtypes of the OS were classified according to the criteria of the WHO (Slayter et al., 1994) and the histologic grade was determined using the grading system established by Kirpensteijn et al. (2002). This information is included in Table 1.

Cell lines

Canine osteosarcoma D17 cells (ATCC No. CRL-6248) were routinely grown in Dulbecco's modified Eagle high glucose medium supplemented with 1% L-glutamine, 10% fetal calf serum, 1% antibioticantimycotic solution (all products purchased from Sigma-Aldrich, St. Louis MO, USA) in a tissue culture incubator at 37°C and 5% CO₂. For pellet preparation, cells were detached using trypsin (Sigma-Aldrich, Vienna, Austria) or scraped, washed with PBS and pelleted by centrifugation. For histological preparation, cell pellets were fixed in 4% formaldehyde for 24 h at 4°C, embedded in Histogel® (Richard-Allan Scientific, Microm International) as specified by the manufacturer,

dehydrated and paraffin-embedded.

Immunohistochemistry

Serial sections of each sample were cut at 3 μ m and mounted on glutaraldehyde-APES-coated slides. After blocking unspecific staining by incubation in 1.5 % normal goat serum for 30 min at room temperature, sections were incubated with primary and secondary antibodies under conditions indicated in Table 2. Immunoreaction was developed with diaminobenzidine chromogen (D5905, Sigma-Aldrich). Nuclear counterstaining was done by hemalumn. Negative controls where the first antibody was substituted by PBS were included with every immunohistochemistry protocol. Positive controls were canine osteosarcoma cells (D17) and feline and canine kidney.

Immunostaining was evaluated by estimating the number of stained tumor cells $(0\%=0, \le 10\%=1, \le 50\%=2, \ge 50\%=3)$ of stained tumor cells) and the staining intensity (zero=0, mild=1, moderate=2, strong=3). Both values were multiplied to receive a total immunostaining score (IS) on one whole tumor section. Tumor center and invasive tumor front (tumor margin) were evaluated separately if available. Two-tailed Fisher's exact test was used calculate statistical significance of the immunoreactivity scores between dog

Table 1. Tumor samples: summary of animal data, tumor subtype and grade.

Case no	Age	Sex	Breed	Tumor localization	Subtype	Grade	Amp/biop/necrop
dog							
1	9y5mo	f/n	mixed	Humerus	osteoblastic	2	amp*1
2	9y5mo	f/n	mixed	Humerus	osteoblastic	2	amp'1
3	12y7mo	m	mixed	Anal sac	chondroblastic	2	biop
4	7y -	f	Leonberger	Tibia	osteoblastic	2	amp
5	11y2mo	m/n	Dachshund	Scapula	osteoblastic	2	amp ¹
6	8y	m/n	mixed	Humerus	fibroblastic	3	amp ¹
7	5y3mo	f/n	Boxer	Radius	fibroblastic	3	amp
8	8y3mo	f	St. Bernard	Radius	osteoblastic	2	amp ¹
9	1y2mo	m	Rhodesian Ridgeback	Ulna	mixed	2	amp
10	14y9mo	f	mixed	Mamma	osteoblastic	2	biop
11	11y3mo	f	Munsterlander	Mandibula	osteoblastic	3	amp ¹
12	7y3mo	m	mixed	Elbow	poorly diff.	3	amp
13	9y	m/n	Dobermann	Femur	osteoblastic	2	amp
14	7y1mo	f	Boxer	Lung/meta	telangiectatic	3	biop/metastasis
15	12y6mo	f	mixed	Tibia	chondroblastic	2	necrop ¹
16	8y7mo	f/n	Am. Staff. Terrier	Humerus	osteoblastic	2	amp
cat							
17	11y11mo	m/n	Europ shorthair	Humerus	osteoblastic	2	amp ¹
18	4y	m/n	Europ shorthair	Tibia	osteoblastic	2	necrop
19	15y	m/n	Europ shorthair	Humerus	fibroblastic	2	amp
20	8y	f/n	Europ shorthair	Skull	chondroblastic	na	biop
21	13y	f/n	Europ shorthair	na	mixed	1	biop
22	10y	f	Europ shorthair	Rib	fibroblastic	1	biop
23	na	na	Europ shorthair	Humerus	mixed	2	necrop
24	6y	f/n	Europ. shorthair	na	osteoblastic	1	biop ¹

F: female; m: male; n: neutered; *: soft part of tumor; ': hard part of tumor; na: not available; amp: amputation; biop: biopsy; necrop: necropsy; 1: metastases at time of diagnosis of primary tumor confirmed.

and cat samples (for on-line calculator use http://www.socscistatistics.com/tests/fisher/Default2.aspx).

Western blotting protocol

Protein lysates were prepared as described elsewhere (Scarlet et al., 2015). The protein concentration was estimated using DC^{TM} Protein Assay (BioRad) according to the manufacterer's recommendations. The soluble, supernatant fraction was stored at -80°C until further analysis.

Protein extracts were separated on 10 % polyacrylamide gels for SDS-PAGE electrophoresis under reducing conditions and transferred to PVDF membrane (GE Healthcare, UK). Membranes were blocked using Western Blocking Reagent (Roche Diagnostics, Germany; dil. 1:10 in TBST) according to the manufacturer's recommendation to avoid unspecific antibody binding. Membranes were incubated with specific primary antibodies followed by secondary antibody incubation as indicated in Table 2. Proteins were visualized using Amersham Western Blotting Analysis System (GE Healthcare, UK). All antibodies were diluted in Western Blocking Reagent/TBST (1:10). Post immunodetection, films and membranes (poststained with Coomassie R-250) were scanned with an Image Scanner III (GE Healthcare Life Sciences) and quantified by densitometry analysis using Quantity One software (version 4.4.0, Bio-Rad). Coomassie protein staining was used as a loading control and for normalization. A cell lysate from the well-known canine osteosarcoma cell line D-17 was analyzed as internal control in western blotting experiments.

Results

Immunohistochemistry

Immunohistochemical scores (IS) and cellular distribution are summarized in Table 3 and Table 4. Immunostaining for ezrin revealed moderate to strong cytoplasmic staining in all canine OS including a lung metastasis. In feline OS, ezrin distribution was also cytoplasmic, however, in contrast to canine OS not all feline tumors were positive for ezrin in the tumor cells and generally expressed a lower IS score. The canine OS cell line (D-17) showed a combination of both cytoplasmic and membrane staining with the anti-ezrin antibody (Figs. 1-8).

The phosphorylated form of pan-ERM proteins (ezrinThr567/radixinThr564/ moesinThr558) was predominantly located in the membrane in the D-17 canine osteosarcoma cell line. In the canine and feline OS tumor samples phospho-ezrin Thr567/radixinThr564/ moesinThr558 signal was also observed in the membranes, additionally, endothelia of blood vessels

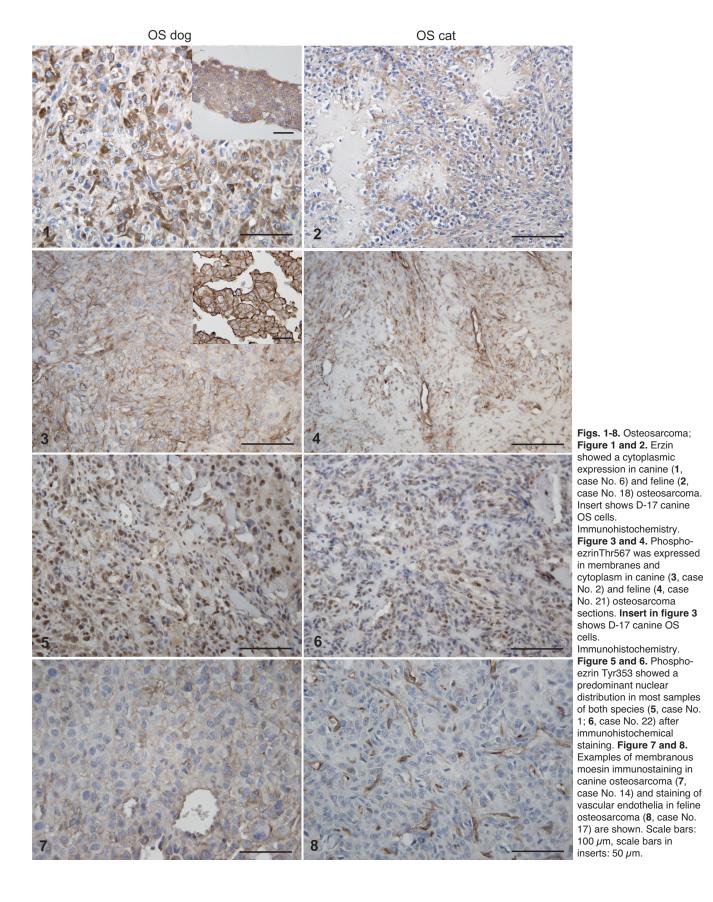
Table 2. Sources, incubation conditions and dilutions of antibodies used in immunohistochemistry and Western blot analyses.

Antibody	Арр.	Incubation time	Temp.	Dilution
Mouse monoclonal anti-ezrin (BD Transduction Laboratories, cat. # 610603)	IHC	ON	4°C	1:250
	WB	ON	4°C	1:500
Rabbit phospho-ezrin (Tyr353) (EnoGene, cat. # E011063, dil. 1:500)	IHC	ON	4°C	1:250
	WB	ON	4°C	1:500
Rabbit monoclonal phospho-ezrin (Thr567)/radixin (Thr564)/moesin (Thr558) (41A3) (Cell Signaling, cat. # 3149)	IHC	ON	4°C	1:250
	WB	ON	4°C	1:500
Rabbit monoclonal anti-moesin (Abcam, cat. # ab52490)	IHC	ON	4°C	1:250
	WB	ON	4°C	1:500
Rabbit polyclonal Akt (Cell Signaling, cat. # 9272)	WB	ON	4°C	1:1000
Rabbit polyclonal phospho-Akt (Ser473) (Cell Signaling, cat. # 9271)	WB	ON	4°C	1:500
Mouse monoclonal GAPDH (GeneTex, cat. # GTX627408)	WB	10 min	RT	1:3000
Anti-mouse BrightVision Poly-HRP (ImmunoLogic, cat. # DPVM110HRP)	IHC	30 min	RT	undil.
Anti-rabbit BrightVision Poly-HRP (ImmunoLogic, cat. # DPVR110HRP)	IHC	30 min	RT	undil.
Amersham ECL-anti-mouse IgG peroxidase-linked species specific whole antibody from sheep (GE Healthcare, cat. # NA931)	WB	30 min	RT	1:5000
Amersham ECL-anti-rabbit IgG peroxidase-linked species specific whole antibody from donkey (GE Healthcare, cat. # NA934)	WB	30 min	RT	1:5000

Table 3. Total immunohistochemical scores (IS) of canine and feline OS tumors.

Score	Dog ezrin	Cat ezrin	Dog p-ezrinThr567	Cat p-ezrinThr567	Dog p-ezrinTyr353	Cat p-ezrinTyr353	Dog moesin	Cat moesin
-	0	1	1	0	1	0	3	4
+	5	6	1	3	1	1	8	3
++	8	0	4	1	7	2	4	1
+++	3	0	10	4	7	5	1	0

^{-:} IS 0; +: IS 1 -3; ++: IS 4-6; +++: IS 7-9. cat ezrin 1 sample n.a.



were strongly labelled. Strongest signal for phosphorylated pan-ERM proteins was present at the tumor margins in both species, whereas for non-phosphorylated ezrin almost no difference between tumor center and tumor margin was determined (Table 4). Most canine samples were found to be in the highest IS; only one canine sample was negative for pan phospho-ERM expression. All feline samples were positive (Figs. 3, 4) with a regular distribution within the IS (Table 3).

The phospho-ezrin form Tyr353 was localized in the nucleus in 13 of 16 canine OS samples, 6 had a cytoplasmic staining, one was negative. In cat 7 of 8 OS samples showed nuclear staining of phospho-ezrin Tyr353. An additional positive immunoreaction was observed in the cytoplasm of tumor cells in sections of both species (Figs. 5, 6). Phospho-ezrin Tyr353 immunostaining was stronger at the tumor margins in 40% of the feline OS samples but only in 15% of canine OS (Table 4).

Expression of moesin in tumor cells was moderate to weak in both species. Explicit membrane staining with anti-moesin was observed in some feline and canine OS including the lung metastasis. Moesin immunostaining clearly marked the endothelia of blood vessels in OS of both species in addition to the tumor cells (Figs. 7, 8). No differences between tumor front and tumor center were distinguishable for moesin expression (Table 4). Statistical evaluation of the low (IS score -/+) and high (IS score ++/+++) immunohistochemical score revealed significant differences (p≤0.05) in ezrin and panphospho-ERM reactivity between dog and cat samples, however the difference in p-ezrinTyr353 and moesin reactivity was not significant statistically. Concerning the primary samples with diagnosed metastasis at time of surgery and samples without confirmed metastasis there was no significant difference in the expression of ezrin or moesin or their phosphorylated forms as assessed by immunohistochemistry. The only lung metastasis that

Table 4. Immunohistochemical distribution of ezrin, phosphorylated ezrin and moesin in canine and feline osteosarcomas.

Antibody	Nuclear positivity		Cytoplasmatic positivity		Membranous positivity		Stronger staining on tumor front	
	cat	dog	cat	dog	cat	dog	cat	dog
ezrin	0/8 (0%)	0/16 (0%)	7/8 (88%)	16/16 (100%)	0/0 (0%)	0/0 (0%)	0/5 (0%)	1/13 (8%)
p-ezrin Thr567	0/8 (0%)	0/16 (0%)	4/8 (50%)	4/16 (25%)	4/8 (50%)	15/16 (94%)	2/5 (40%)	8/13 (62%)
p-ezrin Tyr353	7/8 (88%)	13/16 (81%)	5/8 (63%)	8/16 (50%)	0/8 (0%)	0/16 (0%)	2/5 (40%)	2/13 (15%)
moesin	0/8 (0%)	0/16 (0%)	6/8 (75%)	13/16 (81%)	1/8 (13%)	3/16 (19%)	0/5 (0%)	0/13 (0%)

Tumor front evaluation: only 5 cat samples and 13 dog samples were evaluated due to absence of tumor front on the other samples.

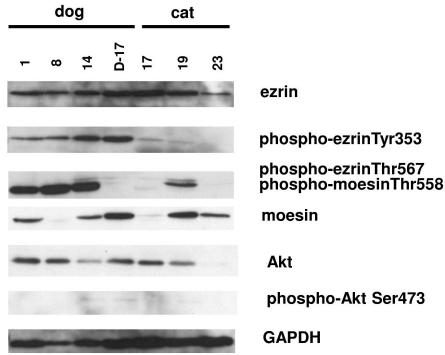


Fig. 9. Western blot detection of the ERM proteins in exemplary dog and cat osteosarcoma samples. Twenty micrograms of protein lysate from different OS subtypes were analyzed as described in Material and Methods section to confirm the specificity of the antibody. Cell lysate from canine osteosarcoma cell line D-17 was analyzed as internal control.

was evaluated showed highest immunohistochemical scores, except for pEzrin Tyr353 where it represented IS 6 (++). Positive controls for immunohistochemistry (feline and canine kidney) reacted as expected, negative controls were blank.

Western blot

Western blot analysis of the osteosarcoma samples revealed similar ezrin expression in all samples of dog and cat OS (Tables 5, 6). However, a moesin band was only present in 82% of dog OS, while it was found in all cat OS in various expression levels. (Table 6, Figs. 9, 10).

Since ezrin phosphorylation status determines its activity in cellular processes, we next performed the analysis using a specific phospho-ezrin (Tyr353) antibody and a pan ERM phospho-ezrin (Thr567)/radixin (Thr564)/moesin (Thr558) antibody. The latter recognizes C-terminal specific threonine phosphorylation in all three forms of ERM proteins; however phosphorylated moesin at Thr558 differs in molecular weight slightly from phosphorylated forms of ezrin and radixin.

To clearly distinguish between signals specific for phospho-ezrin Thr567 and phospho-moesin Thr558, one exemplary cat OS and one dog OS sample were analyzed on a large scale slot, the membrane was cut into several strips and probed with respective antibodies

Table 5. Summary of western blot data.

dog					Thr558		Ser473
1	+++	++	-	++	++++	+++	-
2	++	+	-	+	++++	+	-
3	+++	+/++	+	-	++++	++	+
4	+	-/+	-	+	++++	nd	nd
5	+++	+/++	+	++	++++	+++	+
6	++	+	+	+	++++	+++	-
7	++	+	+	+	++++	nd	nd
8	+++	++	-	+	++++	+++	-
9	+	-	-	-	++++	++	-
10	+/++	+	+	+	++++	-/+	-
11	+	+	-	+	+++	-/+	-
12	++	++/+++	+	+	++++	++	-
13	+	+	+	+	++++	++	-
14	+++	++/+++	+	++	++++	++	+
15	+	+	-	-	++++	++	-
16	++	+/++	+	+/++	++++	nd	nd
cat							
17	+++	+	+	+	-/+	+++	+
18	++	+	+++	+	+++	++	+
19	++	+	++	+++	+++	++/+++	+
20	+/++	-	++	-/+	++	nd	nd
21 +	+/+++	+	++	++	+++	++	+
22 +	+/+++	+	++	++	++	+++	-
23	+/++	-/+	-/+	++	++	+	-
24	++	-/+	+/++	+	+	+++	-

Na: not available; nd: not determined.

(Fig. 11). The bands indicating ezrin, phospho-ezrin Tyr353 and phospho-ezrin Thr567 - upper band - migrated at the same molecular weight at about 80 kDa. The lower band detected with pan phospho-ezrin (Thr567)/radixin (Thr564)/moesin (Thr558) antibody at about 75-78 kDa correlated well with the band specific for moesin and therefore was considered as phosphomoesin Thr558.

Detailed analysis of canine and feline OS samples revealed higher expression of the phosphorylated form of moesin (Thr558) in dog samples as compared to cat OS (Fig. 9, Table 6). On the contrary, the latter samples showed higher expression of the phospho-ezrin Thr567 as compared to dog OS. Substantial differences in phospho-ezrin Tyr353 expression levels were observed between cat and dog OS (Table 6) with higher values in canine OS samples.

As the activation of ERM proteins can lead to the activation of the PI3K-Akt signaling pathway, we also examined Akt and phospho-Akt Ser473 status in these osteosarcoma samples (Fig. 9, Tables 5, 6). Whereas Akt signal was present in all samples with generally similar expression levels for both species, phospho-Akt Ser 473 signal was detected only in 3 out of 13 dog (23% positive) and 4 out of 7 cat (57% positive) OS at very low expression levels.

Western blot results are summarized in Table 5 and 6. No link between the tumor grade or subtype of OS (osteoblastic, fibroblastic, chondroblastic, teleangiectatic, mixed and poorly differentiated) and the amount of ezrin, moesin or the respective phosphorylated forms was found.

Discussion

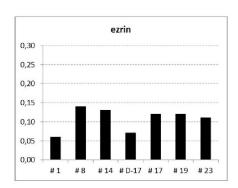
Osteosarcoma is the most common primary bone tumor in dog and cat as well, however dog OS behave differently compared with feline OS. Whereas high incidence of metastases (up to 90%) is described for dogs, only about 10% of feline OS are reported to metastasize (MacEwen, 1990; Dimopoulou et al., 2008). Considering aggressive local bone destruction and metastases occurrence, dog OS closely resembles the

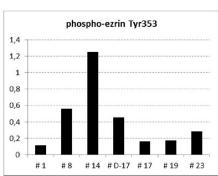
Table 6. Western blot results of ezrin, moesin and Akt expression in dog and cat osteosarcomas.

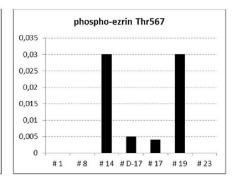
	Dog	Cat
ezrin	16/16 (100%)	8/8 (100%)
p-ezrin Thr567	11/16 (69%)	8/8 (100%)
p-ezrin Tyr353	15/16 (94%)	7/8 (88%)
moesin	13/16 (82%)	8/8 (100%)
p-moesinThr558	16/16 (100%)	8/8 (100%)
Akt *	13/13 (100%)	7/7 (100%)
p-Akt Ser473 *	3/13 (23%)	4/7 (57%)

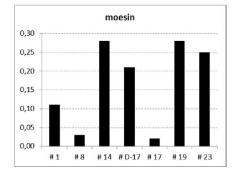
^{*: 13} exemplary dog samples and 7 exemplary cat samples were analyzed.

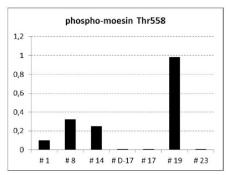
human counterpart (Mueller et al., 2007; Fenger et al., 2014). Pulmonary parenchyma is the most common metastatic site for dog, cat and man (Quigley and Leedale, 1983; MacEwen 1990; Ferrari et al., 2008). The intention of our study was to determine and compare the presence of ERM proteins and their phosphorylation status in the highly aggressive canine OS and the low metastazing feline OS. Tumor metastasis is an important factor affecting survival time and mortality rate. Invasion and metastasis formation is a multistep process including invasion across the basal membrane by the tumor cells, intravasation into blood or lymph vessels, extravasation and growth at the new site (Sahai, 2007). During this process, cells undergo morphological changes driven by the cortical action of the cytoskeleton (Yamaguchi and Condeelis, 2007). Providing a regulated linkage between the plasma membrane and the actin cytoskeleton, the ERM family proteins are involved in determining cell shape, membrane and cell organization, division, migration, and signal transduction (Estecha et al., 2009; Arpin et al., 2011; Neisch and Fehorn, 2011). The metastatic process has been studied extensively in OS and several studies revealed a crucial role of the ERM protein family members in tumor dissemination (Curto and McClatchey, 2004; Fehon et al., 2010). Several processes of how ezrin might influence the metastatic process have been described, however the exact mechanism whereby ezrin contributes to metastasis formation is still not fully elucidated. High ezrin expression in dog and pediatric OS was associated with a significantly shorter median disease-free interval and with early development of metastases (Khanna et al., 2004). Using an osteosarcoma mouse model the same authors have also shown that ezrin is essential for tumor cell survival in the lung after migration from the tumor site and that early metastatic survival was partially dependent on activation of MAPK, but not AKT. However, unexpectedly, not constitutively phospho-

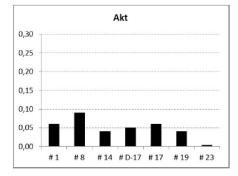












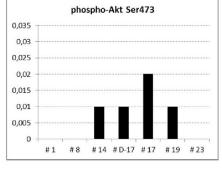


Fig. 10. Semiquantitative densitometrical analysis of proteins detected by Western blot. Protein amount (in arbitrary units) is shown relative to the total protein content loaded per lane.

rylated ezrin expression was sufficient for metastasis formation but only a dynamically regulated ezrin phosphorylation at Thr567 lead to metastatic progression in osteosarcoma (Bitteo et al., 1987). Furthermore, a critical role of ezrin for lung metastasis in OS was also demonstrated by Bulut et al. (2012). Two small molecules (NSC305787 and NSC668394) that inhibit ezrin significantly reduced lung metastases of OS cell lines in a mouse model via inhibition of phospho-ezrin Thr567 levels.

We have carefully analyzed the status of phosphoezrin Thr567 in dog and cat samples using an immunohistochemical as well as a western blot approach. The rabbit monoclonal phospho-ezrin (Thr567)/radixin (Thr564)/moesin (Thr558) (41A3) antibody detects C-terminally phosphorylated forms of all three major ERM proteins and therefore the immunohistochemical analysis cannot distinguish between single phosphorylated proteins. On the contrary, western blot analysis using the antibody mentioned above provides a unique opportunity to discern between phosphorylated ezrin Thr567 and phosphorylated moesin Thr558 by comparing slight differences in the molecular weight of ezrin and moesin, as detected by their specific antibodies with signals obtained using the pan phospho-ERM antibody. Indeed, IHC analysis of all dog and cat samples using pan phospho-ERM antibody revealed positive staining in 15 out of 16 dog and 8 out of 8 cat osteosarcoma samples, respectively. However, there were major differences concerning the subcellular localization of the signal as determined by immuno-

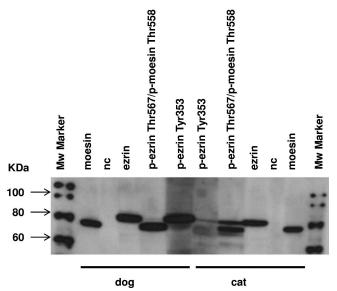


Fig. 11. Size differences between individual ERM proteins as observed in Western blot. The detected MW of ezrin, phospho-ezrin Tyr353 and phospho-ezrin Thr567 (MW ~ 80KDa) differ from the MW of moesin and phospho-moesin Thr558 proteins (MW <80KDa). Representative dog and cat samples are shown.

histochemistry. Membranous staining, indicating the presence of active forms of ERM proteins, was seen in 94% of canine OS whereas only 50% of the feline OS expressed that pattern. Considering the cytoplasmic staining pattern we found 50% of the feline OS but only 25% of the canine OS positive. The numbers of cases in this study is too limited to perform a statistical analysis, so this must be investigated in further studies whether the higher metastatic potential in dog OS is due to the predominating membranous expression of ERM proteins. This would correlate with the findings of Ferrari et al. who reported a significantly longer free survival for patients with only cytoplasmic immunostaining compared to patients with membranous immunostaining in human OS (Ferrari et al., 2008).

Minor discrepancies observed between immunohistochemical and western blot results could be due to the use of different sample parts of the heterogenous OS. Expression of ERM in blood vessels as has also been observed by immunohistochemistry in the present investigation might have an influence on western blotting results. It has been demonstrated that ERM proteins have important differential roles in the thrombin-induced modulation of endothelial cell permeability, with moesin promoting barrier dysfunction and radixin opposing it (Adyshev et al., 2013). It is well known that tumor blood vessels differ significantly from normal blood vessels concerning morphology and permeability (Hashizume et al., 2000), therefore, a closer look at the distribution and function of ERM proteins in tumor blood vessels will be needed.

Subsequent western blot analysis using pan phospho-ERM antibody discovered major differences between the levels of phospho-ezrin Thr567 and phospho-moesin Thr558 expression between dog and cat samples. It was obvious that phospho-moesin Thr558 was more abundant in the OS samples compared to phospho-ezrin Thr567 in both species. This contrast was especially distinct in the dog OS. While all samples were strongly positive for phospho-moesin Thr558, 5 out of 16 OS samples were negative for phospho-ezrin Thr567, whereas the remaining samples revealed only a weak signal corresponding to low amounts of phosphorylated protein. Interestingly, even though this trend was detected also in cat OS, levels of phospho-ezrin Thr567 were higher compared to dog OS and all tested samples were scored as positive. Using the pan ERM antibody in immunohistochemistry it is not possible to discern phospho-moesin Thr558, however, we interpreted the moesin membranous staining as the active/phosphorylated form.

The Akt kinase is known to activate ezrin by phosphorylation on Thr567, and osteosarcoma cells prone to pulmonary metastasis have elevated levels of phospho-Akt (Fukaya et al., 2005; Shiue et al., 2005). Although we have performed analysis of the Akt/p-Akt Ser473 status of the existing dog and cat osteosarcomas, due to low amount of positive samples (4 dogs and 4 cats) we cannot correlate these data to any osteosarcoma

subtype or to a specific pattern of the ERM protein expression.

Estecha and co-workers have shown that moesin is necessary for lung colonization by melanoma cells (Estecha et al., 2009). Moesin silencing markedly reduced the number of invasive cells and the depth of invasion. Interestingly, opposite roles for ezrin and moesin depletion were found in a 3D model of collagen invasion. Only moesin depletion impaired adhesiondependent Rho kinase and subsequent myosin II activation in melanoma cells. As osteosarcoma cells also metastasize to the lung, moesin activation by phosphorylation is an important fact that should also be examined in feline and canine bone tumor. An emerging role of moesin and its phosphorylated form at Thr558 in glioma progression has been described recently (Zhu et al., 2013). Moesin was the major ERM member activated by phosphorylation in human glioblastoma cell lines. Binding to the CD44, a multistructural and multifunctional transmembrane receptor, moesin activated the PI3K-Akt, ERK, and p38MAPK pathways, induced β-catenin nuclear translocation and thereby specifically triggered the activation of β -catenin transcriptional activity. Interestingly, CD44 expression was also found in human osteosarcomas and in OS cell lines as well, and it was positively correlated with lung metastases (Benayahu et al., 2001; Gvozdenovic et al., 2013). Therefore, study of the interaction between moesin and CD44 molecule and its effect on the metastatic potential of dog and feline osteosarcomas remains an attractive topic to be elucidated. However, moesin phosphorylation on Thr558 is not the only known activation of the moesin molecule. G proteincoupled receptor kinases (GRK) phosphorylate moesin at the threonine Thr66 and regulate metastasis in prostate cancer (Chakraborty et al., 2014).

Much less data exist about the function and involvement of phospho-ezrin Tyr 353 in tumor or metastasis progression. In contrast to ezrin Thr567, which switches to the plasma membrane upon activation, ezrin phosphorylated at Tyr353 was found to be translocated into the nucleus in osteosarcomas (Di Cristofano et al., 2010). The exact mechanism of ezrin molecule translocation to the nucleus and its role there is not known yet, although its localization suggests a possible role as a nuclear factor in OS.

Summarizing, we found that cat OS have a higher moesin expression compared to dog OS, however, the active phosphorylated forms of moesin and ezrin Tyr353 were higher in the dog. Although ezrin Thr567 was higher in feline OS, the predominant membranous localization in dog OS samples as observed by immunohistochemistry indicates the presence of the biologically active form. Therefore, it could be speculated that the observed differences in phosphorylated forms of ezrin and moesin status in our pilot study between dog and cat OS could be one of the factors relevant for their different biological behavior. Further knowledge about biological differences of canine and

feline OS could pave the way for novel therapeutic strategies for the treatment of this malignant disease.

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