

Original Article

Relationship of endothelial area with VEGF-A, COX-2, maspin, c-KIT, and DOG-1 immunoreactivity in liposarcomas versus non-lipomatous soft tissue tumors

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Abstract: Soft tissue tumors are rare tumors that show a heterogeneous structure; thus far, their molecular behavior has not been elucidated. The aim of our study was to define the relationship between microvessel density (MVD), evaluated with CD31, and other immunohistochemical markers, such as vascular endothelial growth factor (VEGF), cyclooxygenase-2 (COX-2), CD34, maspin, DOG-1, and c-KIT. Immunostains were done in 55 cases consisting of benign and malignant tumors, such as liposarcomas, dermatofibrosarcomas, and tumors with histiocytic differentiation. Renal tubes were used as external control for VEGF, maspin, and DOG-1. Although DOG-1 is considered a specific marker for gastrointestinal tumors (GISTs), its positivity, correlated with c-KIT and VEGF immunoreexpression, was also shown by dermatofibrosarcomas and tumors with histiocytic and lipomatous differentiation, suggesting its possible pro-angiogenic role. Maspin expression was observed in adipose tissue tumors only. Regarding angiogenesis, 31 of the 55 cases were VEGF-positive, such positivity being directly correlated with COX-2 and CD34 positivity as evaluated in the tumor cells and also with MVD. Although no significant differences in angiogenic activity were found between benign and malignant non-lipomatous tumors, the MVD was directly correlated with the histological type/grade of liposarcomas. Based on these aspects, we conclude that VEGF/COX-2-induced angiogenesis is specific for non-lipomatous tumors, whereas liposarcomas are dependent on the VEGF/maspin angiogenic pathway. The DOG-1/c-KIT/VEGF target may be used for further personalized therapy of soft tissue sarcomas. No data about DOG-1 and maspin positivity in liposarcomas have been published to date.

Keywords: Soft tissue, angiogenesis, maspin, c-KIT, DOG-1, COX-2

Introduction

Soft tissue tumors (STTs) are relatively rare tumors of mesenchymal lineage that can develop in any part of the human body and present a highly heterogeneous microscopic aspect [1, 2]. Although rarely diagnosed, in the case of sarcomas, local recurrences and distant metastases are common, and the mortality rate is higher than 50% [1, 3, 4]. The prognosis and therapy depend on the primary site, size, resection margins, histological subtype, malignancy grade, stage, and molecular signature of the tumor [1, 4, 5]. Moreover, the specific cell origin in each microscopic type of STTs is still unclear [4]; new systems of subclassification have been proposed, the latest introduced by the World Health Organization in 2013 [6]. The biological

behavior of STTs is influenced by some processes, such as angiogenesis, which could inhibit the tumor cell apoptosis [2]. However, other molecular pathways, such as chromosomal translocations, gene fusion [5], and other genomic alterations, have been described. An understanding of these mechanisms would be very useful not only for prognostic assessment but also for targeted therapy of soft tissue sarcomas.

In this study, we aimed to explore the characteristics of some immunohistochemical (IHC) markers whose expressions have not been intensively explored in the literature. Based on previous studies that reported an important prognostic and predictive role of VEGF (vascular endothelial growth factor) -dependent angio-

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Table 1. Main characteristics of antibodies used for the immunohistochemical (IHC) stains (GIST, gastrointestinal stromal tumor)

Antibody (company)	Clone	Dilution	Antigen retrieval	Positive control
Maspin (Novocastra, Newcastle-upon-Tyne, UK)	EAW24	1:25	Incubation with citrate buffer (pH 6.0) - 60 minutes at 100 °C	<ul style="list-style-type: none"> • external - renal tubes • internal - NS
CD31/PECAM-1 (LabVision, Fremont, CA, USA)	JC/70A	1:25	Incubation with high pH-solution (pH 10.0) - 30 minutes at 100 °C	<ul style="list-style-type: none"> • external - placenta • internal - blood vessels - endothelial cells
VEGF-A (LabVision)	VG1	1:50	Incubation with high pH-solution (pH 10.0) - 30 minutes at 100 °C	<ul style="list-style-type: none"> • external - renal tubes • internal - normal mature vessels - endothelial cells
COX-2 (Novocastra)	monoclonal	1:100	Incubation with citrate buffer (pH 6.0) - 60 minutes at 100 °C	<ul style="list-style-type: none"> • external - brain • internal - lymphocytes
CD34 (Dako Glostrup, Denmark)	Qbend/10	1:100	Incubation with high pH-solution (pH 10.0) - 30 minutes at 100 °C	<ul style="list-style-type: none"> • external - placenta • internal - blood vessels - endothelial cells
DOG-1 (Novocastra)	K9	1:50	Incubation with citrate buffer (pH 6.0) - 60 minutes at 100 °C	<ul style="list-style-type: none"> • external - renal tubes • internal - NS
CD117 (Dako)	polyclonal rabbit	1:500	Incubation with high pH-solution (pH 10.0) - 30 minutes at 100 °C	<ul style="list-style-type: none"> • external - intestine • internal - NS

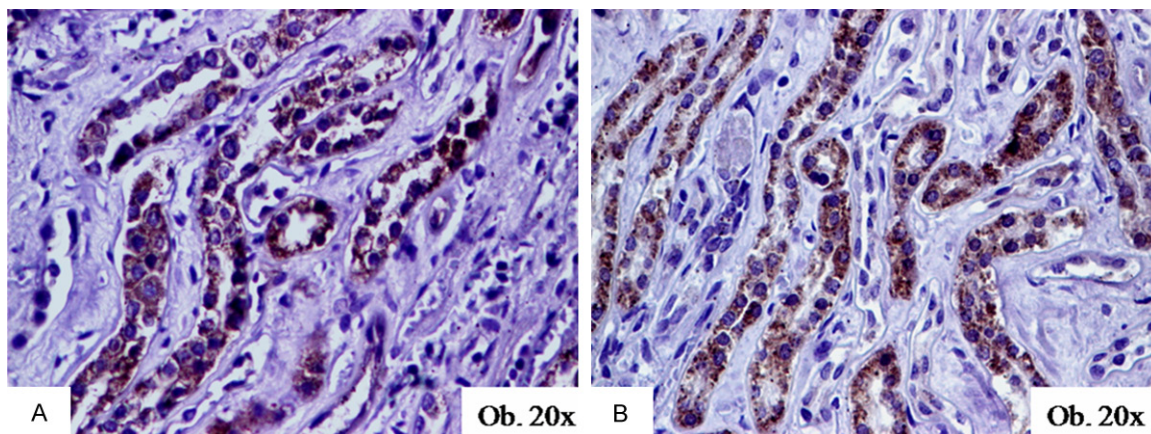


Figure 1. Maspin (A) and DOG-1 positivity (B) in renal tubes that can be used as external positive control.

genesis in STTs [1], we tried to define the particularities by using two markers that have been evaluated in the tumor cells, namely, VEGF-A and cyclooxygenase-2 (COX-2), as well as the endothelial marker CD31.

As original elements, on the one hand, the immunoexpression of maspin and its significance in STTs were analyzed and correlated with other markers. Maspin or serpin B5, a serine protease, is considered to play antiproliferative, proapoptotic, and antiangiogenic roles in carcinomas; however, its role in STTs is not yet known. On the other hand, c-KIT and DOG-1 reactivity within tumor cells was examined for its possible diagnostic and predictive relevance. These markers are known to be relatively specific for gastrointestinal tumors (GIST) [7, 8]. Whereas c-KIT immunoexpression has been observed in several other STTs, such as Kaposi sarcoma [9], tumors with myofibroblastic differentiation, and even melanomas, no extensive studies about DOG-1 specificity have been published thus far. Knowing the predictive value of c-KIT positivity, correlated with the c-KIT mutations in exons 9 and 11 [8], this study aimed to identify those STTs that could respond to receptor tyrosine kinase inhibitors. To the best of our knowledge, no studies about DOG-1 immunoexpression in adipose tissue tumors, or even about the relationship between VEGF, CD31, and maspin in STTs, have been published to date.

Material and methods

Patients and tumor samples

The clinicopathological features of tumors located in dermis and subcutaneous tissues were investigated in 55 consecutive cases that

underwent radical tumor resection from 2010 to 2013. The study was conducted at the Department of Pathology of the University of Medicine and Pharmacy of Tirgu-Mures, Romania. Signed written informed consent was obtained from the patient in each case. All cases were primary tumors; no recurrences or metastases were included. In each of the cases, the age and gender of the patients were considered, and the localization and microscopic type of the tumor were analyzed by using the newest classification of soft tissue tumors [6]. According to this system of classification, liposarcomas were grouped as follows [6]: atypical lipomatous tumors (formerly referred to as well-differentiated liposarcomas), dedifferentiated liposarcomas (liposarcomas with varied non-lipomatous components), myxoid liposarcomas (including the round cell variant), and pleomorphic liposarcomas.

Immunohistochemistry

In each of the 55 cases, immunohistochemical (IHC) stains were done on 5- μ m thick formalin-fixed paraffin-embedded tissues by using a panel of seven antibodies. **Table 1** presents the characteristics of these antibodies. The cases were processed with the Novolink Polymer detection system (Novocastra, Newcastle-upon-Tyne, UK) and developed with the DAB (diaminobenzidine) solution (Novocastra). Counterstaining was done with Mayer's hematoxylin (Novocastra). For negative controls, incubation was done with omission of specific antibodies.

The IHC assessment was done with the use of a Nikon 800E optical microscope with digital camera. The cutoff point for CD34, VEGF-A,

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Table 2. Immunohistochemical profile of soft tissue tumors

Tumor histological type (n = 55)	CD31 median value	VEGF		COX-2		CD34		c-KIT		DOG-1		maspin	
		+	-	+	-	+	-	+	-	+	-	+	-
<i>Liposarcoma</i> (n = 34)	8.35±3.45	27	7	5	29	7	27	5	29	5	29	9	25
<i>Atypical lipomatous tumor</i> (n = 10)	5.98±2.41	8	2	1	9	-	10	-	10	-	10	1	9
<i>Myxoid-type</i> (n = 14)	8.30±4.21	11	3	2	12	-	14	3	11	3	11	3	11
<i>Dedifferentiated type</i> (n = 3)	8.88±2.43	1	2	1	2	-	3	-	3	-	3	-	3
<i>Pleomorphic type</i> (n = 7)	9.44±2.02	7	-	1	6	7	-	2	5	2	5	5	2
<i>Dermatofibrosarcoma</i> (n = 6)	8.98±2.56	1	5	6	-	6	-	1	5	1	5	-	6
<i>Malignant fibrous histiocytoma</i> (n = 4)	9.33±3.68	2	2	4	-	-	4	1	3	1	3	-	4
<i>Fibrosarcoma</i> (n = 3)	8.27±2.31	1	2	3	-	-	3	-	3	-	3	-	3
<i>Benign tumors</i> (n = 8)	9.88±5.11	-	8	8	-	2	6	2	6	2	6	1	7

COX-2, maspin, c-KIT (CD117), and DOG-1 cytoplasmic positivity was set at 10%. The renal tubes were used as external control for VEGF positivity; incidental DOG-1 and maspin positivity was also observed in the renal tubes (**Figure 1**).

The microvessel density (MVD) was evaluated through CD31 staining by digital pictures and the image analysis software ImageJ (NIH). The MVD was determined by counting the positive vessels in the highly vascularized areas (hot spots) at 200x high-power fields. The percentage of positive endothelial area versus the total area of the microscopic field was batch-measured. The ulcerated and inflammatory areas were not taken into account in the assessment of angiogenesis. Although CD34 marked the endothelial cells, it was counted only in the tumor cells and was not used to quantify the MVD.

Statistical analysis

The statistical data were analyzed by using the GraphPad InStat 3 statistical software and descriptive analysis. A *P*-value < 0.05 with 95% confidence interval was considered statistically significant. The frequencies and percentages specific for the analyzed parameters, as well as the means and standard deviations, were used for continuous variables. The t test and ANOVA were used for univariate analysis.

Results

Clinicopathological features

The median age of the 55 patients was 57.23±14.88 years (range, 29-87 years), with a male/female ratio of 1:1.1 (26 males and 29

females). Most of the cases (34 of 55) were liposarcomas. Of the remaining 21 cases, 8 were benign tumors (lipomas and fibrohystiocytic tumors) and 13 were malignant non-lipomatous STTs (**Table 2**). Most of the tumors with lipomatous differentiation were diagnosed in the subcutaneous tissue of the abdominal wall, whereas the other tumor types predominantly developed in the lower extremities.

The endothelial area

By using the CD31 antibody, the median endothelial area (EA) was determined to be 9.08±4.05 (range, 3.17-18.71), without significant differences between benign and malignant tumors (*P* = 0.61). In the case of the benign tumors, the median EA was 9.88±5.11 (range, 3.17-15.78), whereas for the malignant tumors, independent of histological type, the median EA was 8.96±3.93, with a range of 3.95-18.91 (**Table 2**).

Regarding the liposarcomas, the median EA was 8.35±3.45 (range, 3.95-17.48). This was correlated with the tumor type, which was significantly smaller (*P* = 0.02) in atypical lipomas compared with the other histological groups (**Table 2**). The EA was also bigger in liposarcomas compared with lipomas, the latter having a median EA of 3.33±1.21 (*P* = 0.01).

Correlation of the endothelial area with other markers

Independent of the tumor type, the EA was directly correlated with the VEGF (*P* < 0.0001), COX-2 (*P* = 0.03), and CD34 expression in the tumor cells (*P* < 0.0001). No correlations were observed between the EA and the other IHC markers, such as maspin (*P* = 0.11), c-KIT (*P* = 0.09), and DOG-1 (*P* = 0.44).

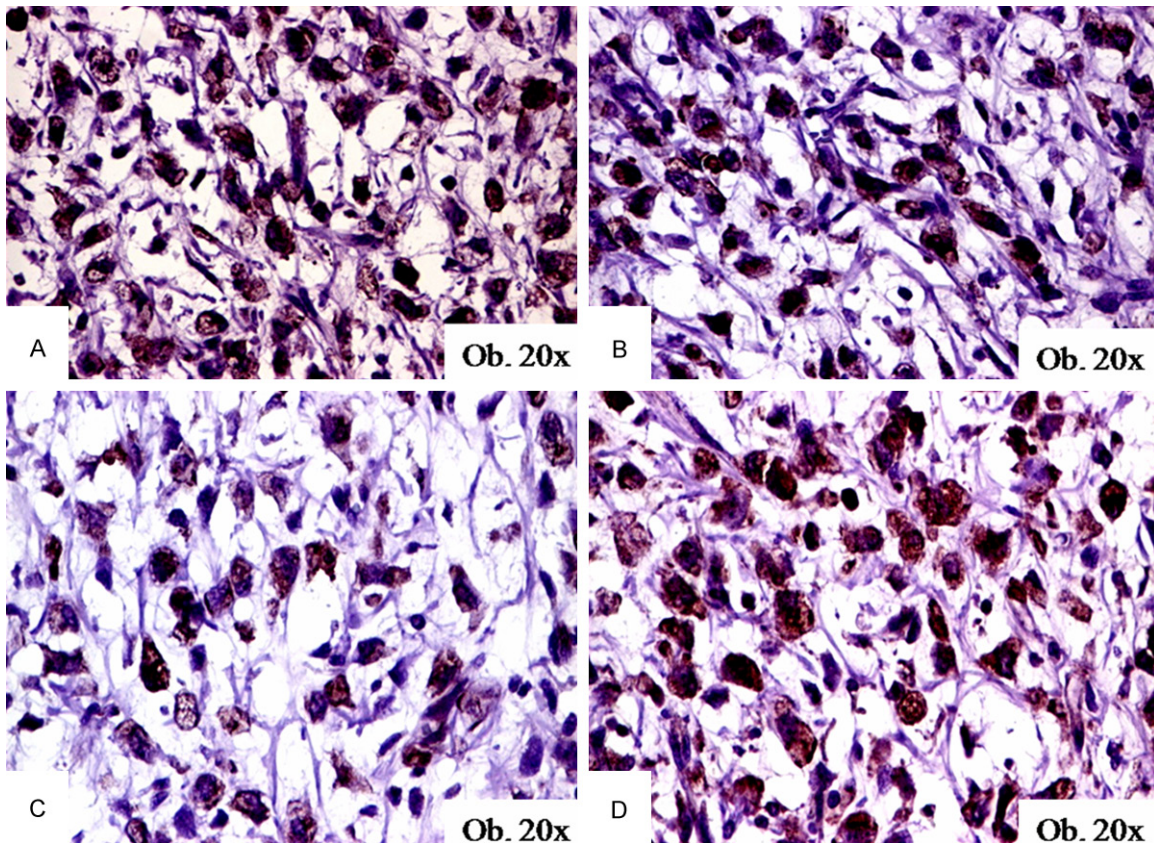


Figure 2. Immunohistochemical expression of DOG-1 (A), maspin (B), c-KIT (C), and VEGF (D) in myxoid liposarcomas.

Immunoexpression of VEGF, COX-2, and CD34 in the tumor cells

Of the 55 cases, 31 showed VEGF positivity in the tumor cells; all of these 31 cases were malignant STTs (Figures 2, 3). COX-2 marked the tumor cells of the 13 malignant non-lipomatous STTs, all benign tumors being COX-2 negative. Of the 34 liposarcomas, only 5 cases showed COX-2 positivity, without any relation to the histological type/grade of differentiation (Table 2).

The CD34 positivity of the tumor cells was observed in 15 of the 55 cases. The two benign CD34-positive cases were lipomas (Table 2). All of the seven pleomorphic liposarcomas were diffusely marked by both CD34 and VEGF (Figure 3).

Immunoexpression of c-KIT and DOG-1 in the tumor cells

DOG-1 positivity was observed in 9 of the 55 cases, both in benign (one lipoma and one benign fibro-histiocytic tumor) and malignant tumors (three myxoid liposarcomas, two pleo-

morphic liposarcomas, one dermatofibrosarcoma, and one malignant fibrous histiocytoma). All of the nine DOG-1-positive cases also showed c-KIT and VEGF positivity (Figure 2). Regarding the DOG-1-maspin correlation, the three non-lipomatous tumors were maspin-negative cases, while the six tumors with lipomatous differentiation also showed maspin positivity (Table 2). DOG-1 positivity was also observed in the mature adipocytes.

Particularities of maspin immunoexpression in the tumor cells

There was marked maspin cytoplasmic positivity in 10 of the 55 cases, all of which presented lipomatous differentiation: one lipoma, one atypical lipomatous tumor, three myxoid liposarcomas (Figure 2) and five pleomorphic liposarcomas were maspin-positive (Table 2).

Discussion

Although the angiogenesis of STTs has been relatively extensively studied in the literature, most reports are focused on sarcomas. Similar to other solid tumors, VEGF is known to be

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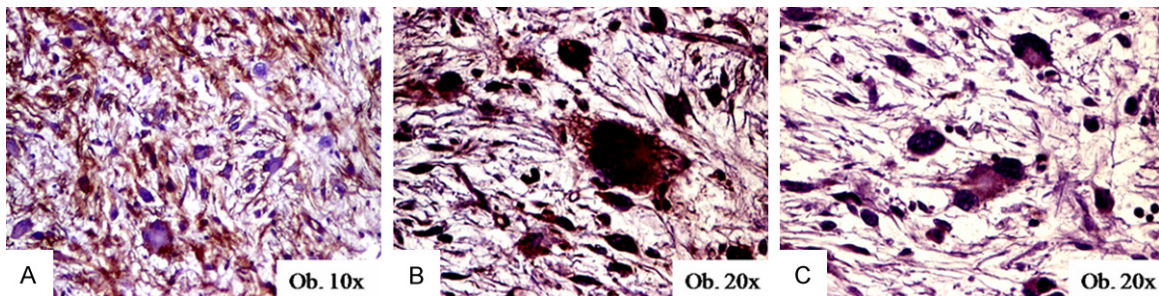


Figure 3. Immunohistochemical expression of CD34 (A), VEGF (B), and DOG-1 (C) in pleomorphic liposarcomas.

secreted by mesenchymal tumor cells and peritumoral mast cells and to induce angiogenesis [10]. At the same time, VEGF and eight other hypoxia-related genes (MIF, SCD1, P4HA1, ENO1/MBP1, FAM162A/HGTD-P, SLC16A1/MCT1, FN1, and STAT1) seem to inhibit the proapoptotic activity of the *bax* gene, assuming that anti-VEGF therapy could increase not only the rate of apoptosis of endothelial cells but also the rate of induced death of tumor cells [2, 5]. In line with these data, a direct correlation between endothelial area, VEGF, and COX-2 was observed in our study. Because anti-VEGF therapy does not achieve the best results in soft tissue sarcomas, a combined anti-COX-2/anti-VEGF therapy may improve the therapeutic results.

Experimentally, the inhibition of maspin in carcinoma cell lines has been proven to have anti-proliferative and antiangiogenic effects, suggesting that anti-maspin drugs could be synthesized for antiangiogenic purposes. However, the confinement of maspin positivity to adipose tissue tumors suggests that it is not involved in the biological behavior of dermatofibrosarcomas or tumors with histiocytic differentiation. In non-lipomatous tumors, angiogenesis seems to be rather dependent on the VEGF and COX-2 activity, whereas a maspin-dependent pathway could be involved in the histogenesis of liposarcomas. COX-2 overexpression induces angiogenesis, enhancing tumor proliferation, and promotes suppression of apoptosis, which is a negative prognostic factor in soft tissue sarcomas [11].

Regarding DOG-1 positivity, although this biomarker is considered to have higher sensitivity and 100% specificity in distinguishing GISTs from other tumors, including leiomyomas and leiomyosarcomas [7, 8, 12], this study proved

for the first time that DOG-1 can also mark other tumors, such as dermatofibrosarcomas, histiocytomas, and tumors with lipomatous differentiation. Its previously described positivity in salivary gland tumors [13], pseudopapillary neoplasm and acinic cell carcinomas of the pancreas [14, 15], melanomas [15], carcinomas of the endometrium and colorectal segments [15], schwannomas [15], and uterine leiomyosarcomas [16], confirms our supposition. Referring to normal cells, DOG-1 has been previously reported to mark pancreatic centroacinar cells [15], its concomitant expression with chromogranin and synaptophysin suggesting a potential DOG-1-induced neuroendocrine activity [17]. Its expression in mature adipose cells and renal collecting tubules, observed in our study, has not yet been reported in the literature.

The DOG-1/VEGF concomitant expression in the tumor cells of soft tissue sarcomas and in the renal tubes could indicate a possible proangiogenic role of DOG-1. This unreported property of DOG-1 could explain the response of both metastatic renal cell carcinomas and metastatic soft tissue sarcomas to pazopanib, one of the most recently approved angiogenesis inhibitors targeting VEGF-, PDGF-, and c-KIT receptors, which is used for second-line therapy of renal cell carcinoma and soft tissue sarcomas [18].

Conclusions

In STTs with histiocytic differentiation and dermatofibrosarcomas the angiogenic activity is dependent on the VEGF-A/COX-2 pathway, whereas a VEGF-A/maspin relationship seems to be more characteristic of liposarcomas. Combined anti-COX-2/anti-VEGF drugs could be effective for non-lipomatous malignant STTs,

whereas liposarcomas could be responsive to anti-maspin/anti-VEGF therapy. Moreover, concomitant DOG-1/c-KIT/VEGF-A positivity might be a potential useful target for further personalized antiangiogenic/anti-tyrosine kinase therapy of STTs; this suggests a potential role of DOG-1 in tumor angiogenesis. Renal tubes can be used as external control for both DOG-1 and maspin.

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Disclosure of conflict of interest

None.

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