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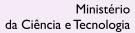
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# Prevalence of vascular-endothelial growth factor, matrix metalloproteinases and tissue inhibitors of metalloproteinases in primary breast cancer

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#### **Abstract**

Our objective was to determine the presence of vascular endothelial growth factor (VEGF), matrix metalloproteinase-2 (MMP-2) and MMP-9 and specific tissue inhibitors of matrix metalloproteinase (TIMP-1 and TIMP-2) in tumor samples obtained from patients with primary breast cancer. We attempted to correlate these findings with the status of the sentinel lymph node (SLN) and clinical-pathological characteristics such as age, tumor size, histological type, histological grade, and vascular invasion. Tumor samples from 88 patients with primary breast cancer were analyzed. The immunoreactivity of VEGF, MMP-2, MMP-9, TIMP-1, and TIMP-2 in tumors was correlated with clinical and pathological features, as well as SLN status. Nonparametric, Mann-Whittney, Kruskal-Wallis, and Spearmann tests were used. Categorical variables were analyzed by the Pearson test. No statistically significant correlation was found between the amount of VEGF, MMP-2, MMP-9, TIMP-1, and TIMP-2 and the presence of tumor cells in the SLN. However, larger tumor diameter (P < 0.01) and the presence of vascular invasion (P < 0.01) were correlated positively with a positive SLN. A significant correlation of higher VEGF levels (P = 0.04) and lower TIMP-1 levels (P = 0.04) with ductal histology was also observed. Furthermore, lower TIMP-2 levels showed a statistically significant correlation with younger age (<50 years) and larger tumor diameter (2.0-5.0 cm). A positive SLN correlated significantly with a larger tumor diameter and the presence of vascular invasion. Higher VEGF and lower TIMP-1 levels were observed in patients with ductal tumors, while higher TIMP-1 levels were observed in lobular tumors.

Key words: Breast cancer; Vascular endothelial growth factor; Matrix metalloproteinase; Metalloproteinase tissue inhibitors; Sentinel lymph node

## Introduction

Pioneering work done by Judah Folkman in the 1970's suggested that tumor angiogenesis was a critical event for cancer growth (1). The interaction between tumor cells and stroma is a vital condition for the beginning of this process, involving molecules that include growth factors, adhesion molecules, angiogenic stimuli, clotting factors, and proteinases (2-4).

Some angiogenic proteins have been described. These include vascular endothelial growth factor (VEGF), basic fibroblast growth factor, tumor necrosis factor alpha, interleukin 8, and matrix metalloproteinases (MMPs) (5).

Endogenous angiogenesis inhibitors such as tissue inhibitors of matrix metalloproteinases (TIMPs) are necessary to block the mitogenic stimuli in the vascular endothelium (5,6). Modifications in the expression of tumor angiogenic factors as well as their inhibitors have been described in various types of cancer, such as colorectal, lung, head and neck and breast cancer, and malignant melanoma (7,8).

VEGF belongs to a family of glycoproteins that includes VEGF-A, VEGF-B, VEGF-C, VEGF-D, and placental growth factor. The main mediator of tumor angiogenesis is VEGF-A, commonly referred to as VEGF (2,8,9). A potent and

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selective mitogenic factor for the endothelium, it elicits a rapid and complete angiogenic response and increased capillary permeability. Produced and secreted by a number of normal cells, it shows marked expression in tumor cells, including breast cancer (10.11).

Several factors contribute to the increased expression of VEGF in breast cancer. Alow intracellular oxygen concentration blocks the degradation of hypoxia-inducing factor- $1\alpha$ , raising its levels and determining intracellular hypoxia, which in turn determines the increase in angiogenic activity for the activation of gene transcription of VEGF (2,12,13). The cleavage of proteins by extracellular MMPs also frees VEGF and endostatin by the cleavage of collagen IV, the latter being an inhibitor of angiogenesis (6,8).

The expansion of solid tumors determines the formation of hypoxic regions, leading to increased production of VEGF, which, in addition to determining the growth and survival of endothelial cells, also acts on some tumor cells, protecting them from apoptosis (10,14,15).

The MMPs are neutral endopeptidases, the family of zinc-dependent proteinases. They play an important role in physiological processes, keeping a balance with the TIMPs. They are also involved in pathological conditions where there is a disruption of this balance, causing diseases such as arthritis, diabetic retinopathy, psoriasis, and cancer (5,7).

Several studies suggest a significant involvement of the MMPs in the process of tumor growth and invasion. A high expression of MMPs, especially MMP-2 and -9, is associated with the histological grade, stage of breast cancer and with the risk of recurrence and death (3,5,7,16,17).

The MMPs are produced by a wide variety of connective tissue cells and also by a variety of malignant cells. Most are in latent forms (zymogens) that are cleaved proteolytically and activated in the extracellular space (5,18).

The MMPs degrade the basement membrane and most components of the extracellular matrix, such as collagen, laminin, fibronectin, and elastin, which are barriers to the normal migration of cells and vascular invasion (6,18).

Molecules with stromal action such as cyclooxygenase-2, adhesion molecules, MMPs, and VEGF accelerate the process of angiogenesis. However, for metastasis to occur, there must be a break-up of the basement membrane, degradation of the extracellular matrix, proliferation and migration of endothelial cells, formation of the vascular lumen, and functional maturation (2,18).

Factors like age, tumor diameter, number of positive axillary lymph nodes, and hormonal and human epidermal growth factor receptor 2 (HER-2) status are currently used in breast cancer to guide the selection of the appropriate adjuvant therapy for women with locoregional disease (19). However, treatment decisions in patients with clinically negative axillary lymph nodes represent a special therapeutic challenge.

Immunohistochemistry has been used in an attempt

to identify tumors with expression of certain molecules that may serve as biological target for new drugs, and can be combated by more specific and effective therapeutic regimens.

For the reasons given above, we determined by immunohistochemistry the expression of angiogenic factors (VEGF, MMP-2 and MMP-9) as well as TIMP-1 and TIMP-2 in tumor samples obtained from patients with primary breast cancer.

Our objective was to correlate these findings with the status of the sentinel lymph node (SLN) and clinical-pathological characteristics such as age, tumor size, histological type, histological grade, and vascular invasion.

#### **Patients and Methods**

Ninety-five patients with a diagnosis of breast cancer, with clinically negative axillary lymph nodes and without distant metastases, were operated at the Academic Hospital in Porto Alegre (HCPA), Federal University of Rio Grande do Sul (UFRGS), between January and December 2004. Tissue samples obtained from the primary tumor and from the SLN were analyzed. This transverse study analyzed the expression of VEGF, MMP-2, MMP-9, TIMP-1, and TIMP-2 in formalin-fixed and paraffin blocks with breast tumor samples. Hematoxylin-eosin preparations were checked and a paraffin block from each case was separated for immunohistochemical studies. Two cases of in situ carcinoma were excluded from the series, and five cases were discarded due to insufficient material, three of them showing a positive SLN for the presence of malignancy. Thus, 88 patients were included in the final analysis.

The clinical and pathological features were classified according to the following parameters: 1) presence (+) or absence (-) of tumor involvement in the SLN. The SLN was considered to be positive when it presented a cluster of malignant cells ≥0.2 mm; 2) age, <50 and ≥50 years; 3) tumor diameter, categorized as ≤2.0 cm or between 2.0 and ≤5.0 cm; 4) histological types classified as ductal or lobular; 5) histological grade, classified as grade 1, grade 2, or grade 3, according to the Scarff Bloom Richardson score (20); 6) presence of at least one nest of tumor cells clearly visible inside the vascular lumen (yes) or absence (no) of vascular invasion detected by hematoxylin-eosin.

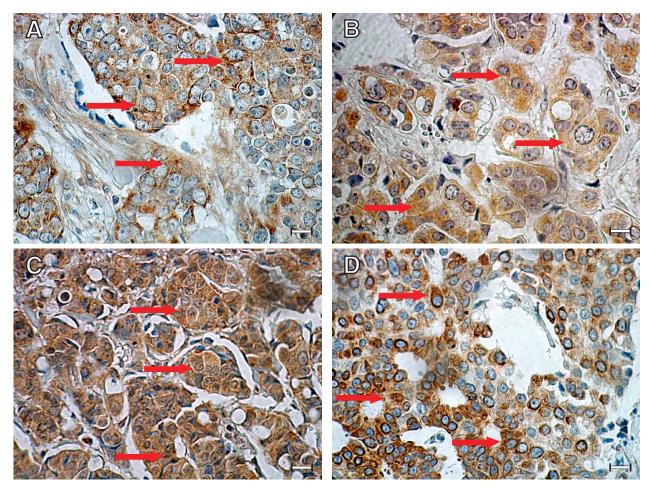
The immunohistochemical technique using the standard avidin-biotin-peroxidase complex was employed to measure the expression of antigens in tumors. The immunoreaction was analyzed in a qualitative and quantitative way. The qualitative estimation was based on the staining intensity of the cytoplasm of neoplastic cells using the following scores: 0 = absence of staining, 1 = slightly positive, 2 = moderately positive, 3 = strongly positive. Protein immunohistochemical expression was quantitated as percentage from 0 (zero) to 100%. The final expression of each case was obtained by the following algorithm: HSCORE =  $\Sigma[(I+1)] \times PC$ , where I

and PC represent the staining intensity and the percentage of stained cells, respectively (21,22).

In order to measure the expression of VEGF we used the polyclonal antibody of rabbit antifactor of vascular human endothelial growth of Signet™, USA, and ultra streptavidin detection at 1:30 dilution, using a sample of breast tumor as external control. The expression of MMP-2 and MMP-9 was measured using the rat monoclonal antibody NCL-MMP2-507, clone 17B11, and the rat monoclonal antibody NCL-MMP9-439, clone 15W2, from Novocastra Laboratories (UK) at 1:40 dilution, using normal hepatic tissue as the external control. The expression of TIMP-1 (tissue inhibitor of MMP-9) was analyzed with the rat monoclonal antibody NCL-TIMP1-485, clone 6F6a (Novocastra Laboratories) at 1:75 dilution, with normal colon tissue as the external control. To determine the expression of TIMP-2 (tissue inhibitor of MMP-2) the rat monoclonal antibody NCL-TIMP2, clone 3A4 (Novocastra Laboratories) was used at 1:25 dilution,

with placental tissue as external control.

Three-micrometer thick paraffin sections were obtained. placed on histological slides treated with poly-D-lysine, deparaffinized with xylol, and rehydrated with 100% alcohol. The slides were washed in tap water and distilled water and incubated in PBS three times for 5 min each time to complete the blockade of endogenous peroxidase. Antigen retrieval was performed with citrate buffer, pH 6, in a microwave oven for 20 min at 90°C. After 15 min at room temperature, the slides were washed again in running and distilled water, and incubated in PBS. After application of 5% hydrogen peroxide, they were washed again and incubated in PBS. The sections were circled with a "Dako-Pen", the primary antibody was added, and the material was left in a moist chamber (Easy Path®, Erviegas, Brazil) in the refrigerator overnight. The slides were then removed from the refrigerator and washed with distilled water. After receiving the secondary labeled streptavidin biotin (LSAB)



**Figure 1.** Immunohistochemical expression of matrix metalloproteinases MMP-2 (A), MMP-9 (B) and tissue inhibitors of metalloproteases [TIMP-2 (C) and TIMP-1 (D)] in neoplastic cells of a breast tumor. The arrows show the immunoreaction in the cytoplasm of the neoplastic cells. Breast cancer cells are shown at 400X magnification (bar = 20 μm in all panels).

antibody (Avidin Biotin, bottle 1, yellow, Dako, USA), the slides were placed in a moist chamber for 30 min. After incubation in PBS, the procedure was repeated by adding the LSAB antibody (streptavidin peroxidase, Dako). The reaction was developed with the chromogenic substrate diaminobenzidine (Dako) for the sections to remain dark (± 30 s) and then were counter-stained with hematoxylin for 20-30 s. After rinsing under running water and application of 2% ammonia, the slides were rinsed in running water again, processed with 100% and mounted with xylol.

Semi-quantitative analysis was carried out by two independent pathologists in the Department of Pathology of HCPA, using an Olympus light microscope, UK, model BX-2 with 10 and 40X objectives.

Qualitative immunohistochemical expression was considered to be positive when the tumor showed at least one cell with cytoplasmic clear and unambiguous brownish staining for each marker utilized, regardless of the intensity of coloration (Figure 1). The quantitative expression of each marker was taken as the average percentage of stained cells in at least 10 microscopic fields.

Statistical analysis to compare the association between SLN-positive groups versus SLN-negative groups and the factors under study and with the other variables selected was performed using the statistical program SPSS V15.

Continuous variables such as age and tumor diameter were compared by the Student t-test. VEGF, MMP-2, -9, TIMP-1, and -2 were correlated with the other clinical-pathological parameters by the non-parametric Mann-Whitney U-test, and the correlation with the histological grade was calculated by the non-parametric Kruskal-Wallis test. The Pearson chisquare test was used to analyze the categorical variables. The correlation of angiogenic factors with one another was calculated by the Spearman correlation coefficient. The confidence interval was set at 95% (95%CI) and values of P < 0.05 were considered to be statistically significant.

The project was approved by the Postgraduate Research Group and Medical Ethics Commission of HCPA (#01/88CNS).

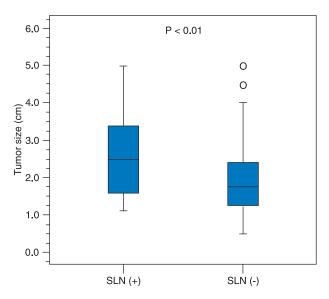
### Results

Table 1 shows the distribution of clinical-pathological and immunohistochemical data according to SLN (+) and SLN (-) status. Mean patient age of 88 patients was 59 years (range: 33-81 years) and mean tumor diameter was 2.3 cm (range: 0.5-5.0 cm). The most frequent histological type was infiltrating ductal carcinoma with 83 (94%) cases, and infiltrating lobular carcinoma was present in 5 cases (6%). The distribution of histological grade according to

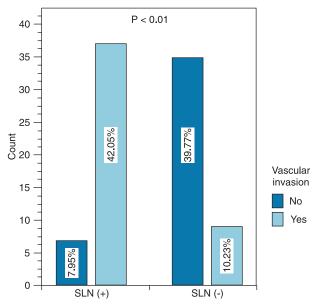
**Table 1.** Clinical-pathological and immunohistochemical data in relation to sentinel lymph node status (SLN): positive and negative SLN groups.

Variables	All (N = 88)	Positive SLN (N = 44)	Negative SLN (N = 44)	
Age <sup>a</sup> (years)	58.7 ± 11.3	58.7 ± 11.8	58.8 ± 11.0	
Tumor diameter <sup>a*</sup> (cm)	2.3 ± 1.0	2.7 ± 1.1	1.9 ± 1.0	
Histological type <sup>b</sup> , N (%)				
Ductal	83 (94%)	42 (96%)	41 (93%)	
Lobular	5 (6%)	2 (5%)	3 (7%)	
Histological grade <sup>b</sup> , N (%)				
G1	23 (26%)	9 (21%)	14 (32%)	
G2	53 (60%)	30 (68%)	23 (52%)	
G3	12 (14%)	5.0 (11%)	7.0 (16%)	
Vascular invasion <sup>b*</sup> , N (%)				
Yes	46 (52%)	37 (84%)	9 (21%)	
No	42 (48%)	7 (16%)	35 (80%)	
VEGF <sup>c</sup>	190 (180-255)	190 (177-239)	190 (180-270)	
MMP-2 <sup>c</sup>	55 (6-204)	63 (4-176)	52 (9-228)	
MMP-9 <sup>c</sup>	180 (100-230)	165 (34-255)	180 (153-196)	
TIMP-1 <sup>c</sup>	368 (340-380)	360 (293-380)	372 (340-387)	
TIMP-2 <sup>c</sup>	249 (170-285)	225 (160-277)	260 (182-322)	
MMP-2/TIMP-2 <sup>c</sup>	0.30 (0.03-0.81)	0.34 (0.01-0.77)	0.26 (0.05-0.88)	
MMP-9/TIMP-1 <sup>c</sup>	0.47 (0.29-0.65)	0.47 (0.10-0.70)	0.49 (0.40-0.60)	

Data are reported as means  $\pm$  SD, score (percent), or median (interquartile range: P25 to P75). The expression of each protein measured by immunohistochemistry is reported as the mean of the algorithm: HSCORE =  $\Sigma[(I + 1)] \times PC$ , where I and PC represent the staining intensity and the percent of stained cells, respectively. VEGF = vascular endothelium growth factor; MMP = matrix metalloproteinase; TIMP = tissue inhibitors of matrix metalloproteinase. \*P < 0.05, significant correlation with positive SLN (at-test, bchi-square test, cMann-Whitney U-test).



**Figure 2.** Correlation between tumor diameter and sentinel lymph node status (SLN): positive (+) and negative (-) SLN. P < 0.01 (Spearman).



**Figure 3.** Correlation between the presence of vascular invasion and sentinel lymph node status (SLN): positive (+) and negative (-) SLN. A positive correlation was observed between the presence of vascular invasion and the occurrence of a positive SLN; P < 0.01 (Spearman).

Table 2. Attempts to correlate the angiogenic factors VEGF, MMP-2 and MMP-9 with SLN, age, tumor diameter, histology, and vascular invasion.

Variables	N (%)	VEGF	MMP-2	MMP-9
SLN				
Positive	44 (50%)	190 (177-239)	63 (4.0-176)	165 (34-255)
Negative	44 (50%)	190 (180-270)	52 (9.0-228)	180 (153-196)
Age (years)				
<50	21 (24%)	190 (178-279)	40 (1.0-175)	160 (43-190)
≥50	67 (76%)	190 (180-225)	80 (3.0-190)	180 (120-255)
Tumor diameter (cm)				
≤2	47 (53%)	190 (180-270)	40 (6.0-210)	180 (120-240)
>2 and ≤5	41 (47%)	190 (176-248)	75 (7.0-210)	170 (53-207)
Histological type				
Ductal	83 (94%)	190 (180-264)*	54 (6.0-210)	180 (100-219)
Lobular	5.0 (6%)	170 (158-185)	84 (23-267)	186 (50-284)
Histological grade				
G1	23 (26%)	186 (176-210)	126 (6.0-186)	180 (46-196)
G2	53 (60%)	190 (180-255)	56 (10-240)	170 (98-217)
G3	12 (14%)	190 (182-277)	11 (0.0-209)	180 (118-262)
Vascular invasion				
Yes	46 (52%)	190 (176-270)	73 (6.0-169)	180 (57-255)
No	42 (48%)	187 (180-255)	45 (8.0-236)	180 (128-190)

Correlation coefficients were calculated by the Spearman correlation method. Data are reported as score (percent) or median (interquartile range: P25 to P75). The expression of each protein measured by immunohistochemistry is reported as the mean of the algorithm: HSCORE =  $\Sigma[(I+1)]$  x PC, where I and PC represent the staining intensity and the percent of stained cells, respectively. SLN = sentinel lymph node; VEGF = vascular endothelial growth factor; MMP = matrix metalloproteinase. \*P < 0.05, ductal compared to lobular histological type (Mann-Whitney U-test; Kruskal-Wallis H-test for histological grade).

**Table 3.** Attempts to correlate the specific inhibitors TIMP-1, TIMP-2, MMP-2/TIMP-2 and MMP-9/TIMP-1 with SLN, age, tumor diameter, histology, and vascular invasion.

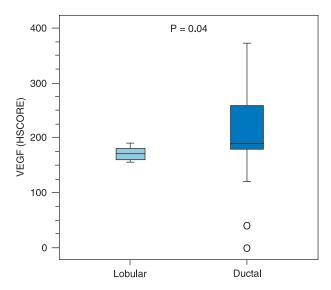
Variables	N (%)	TIMP-1	TIMP-2	MMP-2/TMP-2	MMP-9/TIMP-1
SLN					
Positive	44 (50%)	360 (293-380)	225 (160-277)	0.34 (0.01-0.77)	0.47 (0.10-0.70)
Negative	44 (50%)	372 (340-387)	260 (182-322)	0.26 (0.05-0.88)	0.49 (0.40-0.60)
Age (years)					
<50	21 (24%)	360 (330-370)	178 (149-237)	0.29 (0.01-0.95)	0.47 (0.11-0.58)
≥50	67 (76%)	372 (340-388)*	264 (190-332)*	0.35 (0.04-0.72)	0.48 (0.31-0.67)
Tumor diameter (cm)					
≤2	47 (53%)	372 (340-380)	270 (190-291)*	0.18 (0.03-0.70)	0.48 (0.31-0.67)
>2 and ≤5	41 (47%)	360 (319-380)	190 (160-270)	0.44 (0.03-0.87)	0.47 (0.14-0.63)
Histological type					
Ductal	83 (94%)	368 (332-380)	234 (170-285)	0.31 (0.03-0.82)	0.47 (0.30-0.63)
Lobular	5 (6%)	388 (360-394)*	270 (255-366)	0.22 (0.11-0.82)	0.48 (0.14-0.72)
Histological grade					
G1	23 (26%)	380 (340-388)	225 (160-285)	0.57 (0.03-0.82)	0.50 (0.12-0.57)
G2	53 (60%)	368 (340-380)	249 (170-306)	0.30 (0.07-0.83)	0.47 (0.28-0.66)
G3	12 (14%)	350 (293-386)	245 (200-281)	0.04 (0.00-0.60)	0.50 (0.33-0.72)
Vascular invasion					
Yes	46 (52%)	360 (296-380)	230 (160-230)	0.32 (0.02-0.79)	0.49 (0.15-0.73)
No	42 (48%)	372 (340-381)	255 (189-334)	0.24 (0.04-0.84)	0.47 (0.33-0.55)

Correlation coefficients were calculated by the Spearman correlation method. Data are reported as score (percent) or median (interquartile range: P25 to P75). The expression of each protein measured by immunohistochemistry is reported as the mean of the algorithm: HSCORE =  $\Sigma[(I+1)]$  x PC, where I and PC represent the staining intensity and the percent of stained cells, respectively. SLN = sentinel lymph node; TIMP = tissue inhibitors of matrix metalloproteinase; MMP = matrix metalloproteinase. \*P < 0.05 (Mann-Whitney U-test; Kruskal-Wallis H-test for histological grade).

the Scarff Bloom Richardson score was classified as G1: 23 (26%), G2: 53 (60%) and G3: 12 (14%). Vascular invasion was documented in 46 (52%) cases. Tumor samples were analyzed in terms of the presence or absence of SLN involvement. Tumor diameter (P < 0.01; Figure 2) and the presence of vascular invasion in the primary tumor were correlated significantly (P < 0.01; Figure 3) with a positive SLN.

Table 1 also shows the immunohistochemical expression of VEGF, MMP-2 and -9 and TIMP-1 and -2 in the cytoplasm of neoplastic cells, measured by the means of HSCORE and the ratio between MMPs and their specific inhibitors (MMP-2/TIMP-2 and MMP-9/TIMP-1). VEGF presented a median score of 190 (180-255); MMP-2 presented a median score of 55 (6-204), with 14 (15.9%) negative cases; MMP-9 presented a median score of 180 (100-230), with 10 (11.4%) negative cases; TIMP-1 presented a median score of 368 (340-380), and TIMP-2 a median score of 249 (170-285). No significant correlation between the immunohistochemical expression of the proteins cited above and SLN status was observed.

Table 2 presents the immunoreactivity of VEGF, MMP-2 and MMP-9 in relation to the SLN status and other variables. Tumors with the lobular histological type showed a significantly lower VEGF than the ductal type (P = 0.04; Figure 4).



**Figure 4.** Correlation between vascular endothelial growth factor (VEGF) levels and histological tumor type. A positive correlation between VEGF expression and histological type was observed (P = 0.04; Spearman).

In Table 3, SLN status and the variables cited above were correlated with TIMP-1 and TIMP-2 levels and with the ratio

**Table 4.** Summary of statistically significant correlations between VEGF, MMP-2, MMP-9, TIMP-1, TIMP-2, MMP-9/TIMP-1 levels identified by the Spearman correlation method.

Factors	VEGF	MMP-2	MMP-9	TIMP-1	TIMP-2	MMP-2/TIMP-2	MMP-9/TIMP-1
VEGF							
rho	-		0.26				0.32
Р	-		0.01				0.02
MMP-2							
rho		-			0.36	0.92	
Р		-			< 0.01	<0.01	
MMP-9							
rho	0.26		-				
Р	0.01		-				<0.01
TIMP-1							
rho				-			
Р				-			
TIMP-2							
rho		0.36			-		
Р		<0.01			-		
MMP-2/TIMP-2							
rho		0.92				-	
Р		<0.01				-	
MMP-9/TIMP-1							
rho	0.32		0.95				-
P	0.02		<0.01				

SLN = sentinel lymph node; VEGF = vascular endothelium growth factor; MMP = matrix metalloproteinase; TIMP = tissue inhibitors of matrix metalloproteinase; rho = Spearman rank correlation. Data are reported only for P < 0.05 (Spearman correlation coefficient).

between MMP-2 and MMP-9 and their specific inhibitors (MMP-2/TIMP-2 and MMP-9/TIMP-1). TIMP-1 levels were higher in tumors of the lobular type (P = 0.04) and TIMP-2 levels were higher in older patients (P < 0.01) and in smaller tumors (P = 0.01). The MMP-2/TIMP-2 and MMP-9/TIMP-1 ratios showed no significant correlation.

The correlation of VEGF, MMP-2, MMP-9, TIMP-1, TIMP-2 levels with the MMP-2/TIMP-2 and MMP-9/ TIMP-2 ratios is shown in Table 4. VEGF levels correlated significantly with the MMP-9 factor (P = 0.01). A positive correlation was also observed between MMP-2 and TIMP-2 levels (P < 0.01).

#### **Discussion**

The recognition that the occurrence of tumor growth and metastasis depends on the angiogenesis process has promoted the investigation of prognostic and predictive factors such as hormone receptor, HER-2, VEGF and MMPs, in order to elucidate the significance of angiogenesis in breast cancer. The analysis of the expression of these biomarkers in tissues of breast cancer can be of great prognostic value in assessing the metastatic potential of tumor and survival

of patients (3,23).

Several studies have suggested a relationship between VEGF, MMP-2, MMP-9, TIMP-1, and TIMP-2 expression in primary breast cancer and disease prognosis. Some of them suggested that patients with high levels of VEGF, MMP-2 and MMP-9 expression in the primary tumor might have a more aggressive clinical course, with presentation of early metastases. However, these results are still conflicting and not clear (13,24,25).

Analyzing the literature, it was observed that the quantification of immunohistochemical expression of proteins in tumors is determined in different manners, using the most varied scores. All authors consider the presence of brown staining in neoplastic cells as a parameter for positivity. Some authors consider only the percentage of stained cells to set expression (26,27), while most link the percentage of cells stained with the intensity of staining, but using various formulas (28-30). We applied the algorithm HSCORE, widely described in the literature (19,20), in an attempt to use a method of compression and easy reproducibility.

In the present study, larger tumor diameter (P < 0.01) and the presence of vascular invasion (P < 0.01) were correlated significantly with the presence of a positive SLN.

When comparing the expression of VEGF, MMP-2, MMP-9 and their specific inhibitors TIMP-1 and TIMP-2 to the presence or absence of tumor cells in the SLN, no statistically significant correlation was found.

Several authors, such as Nakopoulou et al. (31) and Pellikainen et al. (26), found high expressions of MMPs associated with small tumors (TI-2) and negative axillary lymph nodes. However, other authors, such as Kim et al. (32) and Hao et al. (30), found a significant association of VEGF and MMP-9 with advanced histological grade and positive axillary lymph node.

We found a significant correlation between higher VEGF expression (P = 0.04) and lower TIMP-1 levels (P = 0.04) with the ductal histological type. Furthermore, lower TIMP-2 levels also showed a significant correlation with younger age (<50 years) (P = 0.002) and a larger tumor diameter (2.0-5.0 cm; P = 0.01).

Nakopoulou et al. (33) studied 133 cases of invasive breast cancer and detected an increased TIMP-1 expression in 60% of cases, with an inverse correlation between TIMP-1 and histological grade (P = 0.003). The cases with increased TIMP-1 expression were correlated with favorable prognosis factors, mainly with negative axillary lymph nodes. Yet, elevated TIMP-1 levels were associated with better survival (P = 0.01).

Kim et al. (32), when analyzing 63 cases of T1-T2 breast tumors, did not find a correlation of TIMP-2 expression with histological grade or axillary lymph node status. A study of Têtu et al. (34) found the TIMP-2 expression associated with the ductal histological type.

In order to analyze the correlation of the balance between MMPs and their inhibitors with SLN and other variables under study, we calculated the MMP-2/TIMP-2 and MMP-9/TIMP-1 ratios. We did not find a significant correlation between these ratios and SLN involvement.

Jinga et al. (22) analyzed this ratio in benign and malignant breast tissues and found a significant difference in malignant tissues, suggesting that an abnormal balance between MMPs and TIMPs plays an important role in tumor growth patterns.

Analyzing the relationship MMP-2/TIMP-2, Têtu et al. (34) found a worse 5-year survival in those patients with low expression of TIMP-2 and high expression of MMP-2, but did not find an association between the expression of

TIMP-2 and overall survival. In contrast, Nakopoulou et al. (31) observed a better survival in cases with positive TIMP-2 and negative MMP-2 expression, and also in cases that were positive for both factors. Other investigators also reported that changes in the balance between MMPs and TIMPs with an increase in the activity of MMPs played an important role in tumor invasion and metastasis (22,34).

Currently, the angiogenic factors represent an attractive therapeutic target for a new class of anticancer drugs. Although the use of antiangiogenic drugs for the treatment of cancer has increased in recent decades in basic clinical research, their clinical benefits are relatively low (35-38). Bevacizumab (Avastin, Genentech Inc., USA), is the most investigated antiangiogenic agent (2,39). Miller et al. (40) recently published a randomized phase III study comparing the use of paclitaxel associated with bevacizumab versus paclitaxel alone for the initial treatment of metastases in breast cancer. The results showed a significant increase in objective response rates (36.9 vs 21.2%, P < 0.001) and in disease-free survival (median, 11.8 vs 5.9 months, P < 0.001), both in favor of the paclitaxel/bevacizumab combination.

A better definition of the biological nature of tumors allows individual approaches based on knowledge of the molecular profile of each patient. The use of antiangiogenic agents in combination with other chemotherapeutic agents can improve the efficiency of treatment and prolong the lives of patients with metastatic disease and prevent recurrence in those with initial disease (3,10).

The use of angiogenic factors to identify patient prognosis in early breast cancer is not clear. The data presented in the literature are conflicting and this will remain a problem until the mechanisms involved in carcinogenesis and metastatic invasion are better understood.

In our series of patients with primary breast cancer, a larger tumor diameter and the presence of vascular invasion were correlated with a positive SLN. Higher VEGF levels and lower TIMP-1 levels were shown to be more frequently associated with tumors of the ductal type, while higher TIMP-1 levels were found in patients with tumors with lobular type. Lower TIMP-2 levels showed also a significant correlation with younger patient age and larger tumor diameter. These results warrant further studies.

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