# Prognostic phenotypic classification for canine mammary tumors

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Abstract. Mammary neoplasms are a heterogeneous form of disease, and in order to determine its course and biological features with more accuracy, investigations based on tumor phenotypes are required. The aim of the present study was to propose and validate a phenotypic classification for canine mammary tumors and to assess any association between clinicopathological characteristics, survival and prognosis. For the immunohistochemistry analysis, the primary antibodies against estrogen receptor-α, progesterone receptor, human epidermal growth factor receptor 2 (HER-2)/neu and E-cadherin were used. A total of 110 canine mammary tumors were investigated; 42 tumors were classified as luminal A, 41 as luminal B, 17 as triple-negative and 10 as HER-2-positive. The luminal A and B phenotypes were associated with improved prognosis, whereas HER-2positive and triple-negative tumors were more aggressive, and exhibited a significant association with the occurrence of metastasis, a worse Tumor-Node-Metastasis classification and shorter survival time (P<0.05). In addition, there were different levels of E-cadherin expression intensity observed among the four tumor profiles investigated. Luminal A and B phenotypes presented an upregulation of E-cadherin compared with the HER-2 positive and triple-negative

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Abbreviations: ER, estrogen receptor; HER-2, human epidermal growth factor receptor 2; PR, progesterone receptor; TNM, Tumor-Node-Metastasis

Key words: dogs, E-cadherin, mammary tumors, phenotypes, prognostic

phenotypes (P<0.05). From the results of the present study, the proposed immunohistochemical panel and phenotypic classification techniques could be useful diagnostic tools with a good technical applicability in veterinary oncology. The analysis of E-cadherin expression in the panel of tumor markers allowed a more accurate classification for determining the biological features of the mammary tumor.

## Introduction

Mammary tumors are the most common types of cancer affecting unspayed bitches or bitches that are spayed late in life (1-5). This disease represents ~52% of all neoplasms affecting bitches, and 41-53% of mammary tumors are histopathologically diagnosed as malignant (1-7). Studies conducted in Brazil between 2010 and 2012 have demonstrated that the incidence rate of malignant mammary tumors ranges from 60-82% (8,9). Therefore, studying mammary neoplasms is of great importance due to the high rates of mortality and morbidity that affect this animal population (10,11).

Clinical and histological prognostic factors for mammary tumors have been previously studied and have been suggested to be important parameters in determining the possible outcome and progression of the disease (3,12). The most important clinical characteristics are tumor size, lymph node status and the presence of distant metastasis (13). However, they are still incipient and cannot be used as precise and reliable predictive and/or prognostic factors, particularly in the early stages of the disease (10).

Previous histological classifications for canine mammary tumors described in the literature failed to identify potential biological differences among histological subtypes, particularly in patients with the same disease status that have the same anatomical, morphological and grade histological structure (14-18). A number of researchers have suggested new histopathological classifications, or have made attempts to modify the existing ones, resulting in subjectivity during interpretation (14-18). In addition, due to many different

classification systems, oncologists have found it difficult to select the one that best defines and characterizes the disease as well as its progression (10).

Cell phenotypes can be used as a new tool that contributes to the understanding of tumor biology (10), so cancer cell immunostaining should be emphasized as an effective tool in the clinical setting. A number of immunohistochemical markers have been proposed and a panel of markers has been already established for use in the clinical prognosis of women with breast cancer (19-21). Importantly, the analysis of this panel of markers can identify a specific phenotypic tumor profile, which is associated with a particular biological behavior (22,23). This panel includes the estrogen receptor (ER) and progesterone hormone receptor (PR), human epidermal growth factor receptor 2 (HER-2)/neu oncogene, the cell proliferation marker Ki-67 and p53 (10). However, in veterinary medicine, there are few studies currently focusing on this subject and the published results are conflicting (24-26).

Therefore, the objective of molecular research in breast cancer, in addition to defining specific biological profiles, is to identify novel potential therapeutic targets in order to increase the control and the treatment of this disease. Previous findings have provided important prognostic and predictive information, as well as a better understanding of the complex molecular mechanisms underlying tumorigenesis (22,27). As a result, different treatment approaches were optimized and thus, the ability to identify molecular subtypes has become an important element for the management of breast cancer (28).

By combining the protein expression of ERs, PRs and HER-2, three markers widely used in the diagnosis of breast cancer in women, it is possible to identify the luminal groups A, luminal B, HER-2-positive and triple-negative (22). This new form of classification has gained popularity and credit among the scientific community as it has been discovered that the groups are associated with specific treatments (29-31). The luminal group presents a favorable prognosis and responds to hormone therapy, whereas monoclonal therapy is indicated in the HER-2 group (32,33). The targeted therapies against HER-2 positive tumors are very effective, both in the adjuvant and the metastatic setting (34,35). Trastuzumab is a humanized monoclonal antibody that improves response rates, decreases the progression of the disease and improves survival time when used alone or added to chemotherapy in metastatic breast cancer (36). The triple-negative group has worse prognosis and requires specific treatments (37,38). It is expected that patients with this profile will not benefit from the use of trastuzumab or hormonal therapies, which results in challenges when prescribing and administering treatments in patients with triple-negative mammary tumor (39,40).

In addition, E-cadherin is considered an important prognostic immunohistochemical marker. This molecule is a cell adhesion protein and an important regulator of the epithelial phenotype. The low expression of E-cadherin is associated with the malignant progression of the disease (41), and the low expression of this protein occurs in processes involved in the epithelial-to-mesenchymal transition (EMT). EMT promotes the acquisition of mesenchymal characteristics in epithelial cancer cells with the invasion and spreading of cancer

cells (41,42). Canine mammary tumors have been observed to be significantly associated with the loss of E-cadherin expression, resulting in a poor prognosis (43).

Therefore, the purpose of the present study was to define a phenotypic immunohistochemical classification for canine mammary tumors in four groups: i) luminal A (ER<sup>+</sup> and/or PR<sup>+</sup>, and HER-2<sup>-</sup>); ii) luminal B (ER<sup>+</sup> and/or PR<sup>+</sup>, and HER-2<sup>+</sup>); iii) HER-2 positive (ER<sup>-</sup>, PR<sup>-</sup> and HER-2<sup>-</sup>); and iv) triple-negative (ER<sup>-</sup>, PR<sup>-</sup> and HER-2<sup>-</sup>). In addition, the present study investigated E-cadherin expression in these phenotypes in order to examine the association between tumor classification and prognosis also in the canine species.

#### Materials and methods

Experimental group. The experimental group consisted of 110 bitches, enrolled between January 2011 and December 2013, of different breeds and ages (3-15 years). In all dogs examined, the mammary cancer developed spontaneously. All patients were treated in the Obstetric and Reproduction Department of the Veterinary Hospital Governador Laudo Natel of Faculdade de Ciências Agrárias e Veterinárias/Universidade Estadual Paulista, Jaboticabal, Brazil, and the veterinary clinics of São José do Rio Preto (São Paulo, Brazil). The owners of each dog provided written consent for the use of the samples, and the present study was approved by the The Ethics Committee on Animal Experimentation of the Faculty of Medicine of São José do Rio Preto prior to the study start. All dogs were treated surgically, and none of the dogs received any additional anticancer treatment prior to or following mastectomy.

Following excision of the tumor, the patients were followed up for 1-18 months. During the follow-up time, the veterinarians evaluated tumor metastasis and recurrence, as well as the cause of mortality in the animals. For histopathological diagnosis, the tumor biopsies collected were classified according to Goldschmidt *et al* (18).

The parameters employed for the classification of clinical tumor staging were in accordance with the Tumor-Node-Metastasis (TNM) system established by the World Health Organization for canine mammary gland tumors (44.45).

Immunohistochemistry (IHC). The IHC procedure was performed following the method described by Lopes et al (46). Tumor samples were embedded into paraffin blocks and cut into 3- $\mu$ m sections. The samples were prepared on silanized glass slides prior to deparaffinization. The sections were rehydrated in an ascending range of alcohol concentrations and incubated with 3% hydrogen peroxide for 30 min. Antigen retrieval was performed by heating at 95°C in buffer for 35 min. The slides were incubated with bovine serum albumin (BSA). The slides were incubated at 4°C overnight with the primary antibodies (Table I). After being washed with phosphate-buffered saline (PBS) for 15 min, incubation was carried out with Starr Trek Universal HRP Detection kit (Medical Biocare, Concord, CA, USA), consisting of the secondary antibody 'anti-mouse, rabbit and goat immunoglobulin with biotin' for 1 h and 'streptoavidin complex with peroxidase' for 30 min, followed by washes with PBS for

15 min. Subsequently, 0.5% 3,3'-diaminobenzidine tetrahydrochloride was applied to the slides for 2-5 min at 20-22°C. The slides were counterstained with Harris' hematoxylin for 40 min. Negative controls were obtained by omitting the primary antibody.

The slides were scanned using the Panoramic Digital Slide Scanner (magnification, x40; 3DHISTECH®) and Panoramic Viewer software (3DHISTECH®) (47).

Analysis and interpretation of IHC. For the ERs and PRs, the immunoreactivity index of Allred *et al* (48) was applied. The final score was calculated according to the quantity of marked cells and the intensity of the staining. The scores ranged between 0 and 8. Samples with a score of 0-1 were considered negative, and samples with a score of 2-8 were considered positive. The scores and criteria used to quantify the labeled cells were: i) 0, no labeling; ii) 1, labeling in <1% of cells; iii) 2, 1-10%; 3, 11-33%; iv) 4, 34-66%; and 5, 67-100%. The criteria and scores used to determine the intensity of immunostaining were: i) Absent, 0; ii) weak, 1; iii) moderate, 2; and iv) strong, 3. The staining intensity was determined by eye. The sum of these criteria for each sample determined the final score

HER-2/neu exhibited membrane staining in >10% of neoplastic cells, and the intensity of the staining was assessed according to a previously described semi-quantitative method (49). The criteria used for the score were as follows: i) 0, no staining; ii) 1, weak, incomplete membranous staining; iii) 2, moderate, complete membranous staining in at least 10% of tumor cells; and iv) 3, strong membranous staining in at least 10% of tumor cells. The staining intensity was determined by eye. Cases with a score of 0-1 were considered negative, whereas scores of 2-3 were considered to exhibit HER-2 positive.

Expression of E-cadherin was performed using the modified semi-quantitative immunoreactive score scale according to Remmele and Stegner (50). Immunostaining signal of E-cadherin was observed in the cytoplasmic membrane. The method described by Remmele and Stegner takes into account both the proportion of positively stained cells and the intensity of the staining. The criteria used to quantify the labeled cells were: i) 0, no labeling; ii) 1, labeling in ≤10% of cells; iii) 2, 11-50%; iv) 3, 51-80%; and v) 4, 81-100%. The criteria used to determine the intensity of immunostaining were: i) Zero, no staining; ii) '+', weak staining; iii) '++', moderate staining; and iv) '+++', strong staining. The multiplication of the criteria for each sample determined the final score.

Statistical analysis. All statistical analyses were performed using GraphPad Prism (version 5.0; GraphPad Software, Inc.) software. Raw data were initially subjected to descriptive analyses to determine the normal distribution. Any associations between immunostaining intensity of E-cadherin and the mammary cancer phenotypes of the patients within each group were analyzed using the  $\chi^2$  test. The survival curve was constructed using the Kaplan-Meier method and the differences between the curves were evaluated using a log-rank test and hazard function. P<0.05 was considered to indicate a statistically significant result.

#### Results

Phenotype characterization. In the present study, the luminal A phenotype was the most frequently observed (38.2%), followed by luminal B (37.3%), triple-negative (15.4%) and HER-2 positive (9.1%). Fig. 1 presents the phenotypic profiles according to the immunohistochemical markers ER, PR and HER2. The mammary tumor group exhibited the most frequent histopathological classifications: i) Luminal A (29.0% complex mammary carcinoma, 24.0% tubulopapillary mammary carcinoma and 21.0% mixed mammary carcinoma); and ii) luminal B (37.0% complex mammary carcinoma, 19.0% tubulopapillary mammary carcinoma, 15.0% mixed mammary carcinoma and 15.0% solid carcinoma); iii) HER-2 positive (30.0% complex mammary carcinoma, 30% tubulopapillary mammary carcinoma and 20.0% anaplastic mammary carcinoma); and iv) triple-negative (29.4% complex mammary carcinoma, 23.5% mixed mammary carcinoma and 23.5% tubulopapillary mammary carcinoma).

Mammary tumor phenotypes and clinicopathological characteristics. For the luminal A tumor subtype, only 14.3% (6/42) of the animals presented cutaneous (2/42) and pulmonary (2/42) metastasis and recurrence (1/42). Regarding the TNM classification, the minority of mammary tumors were classified with the worst prognosis. Regarding the luminal B subtype, 22.0% (9/41) of the animals presented hepatic (2/41) and pulmonary (5/41) metastasis and recurrence (5/41). For the TNM classification, stages I and II (30/41) were significantly higher in number compared with stages III, IV and V.

Regarding HER-2 positive, 80.0% (8/10) of animals presented cutaneous (1/10) and pulmonary (7/10) metastasis and 60.0% (6/10) exhibited stage III, IV or V TNM classification. In the triple-negative tumor phenotype, 76.4% (6/10) of animals presented hepatic (1/17) and pulmonary (12/17) metastasis and 53.0% (9/17) of tumors exhibited stage III, IV and V TNM classification.

Table II presents the distribution of frequency (number of cases) of clinicopathological characteristics in the phenotypes luminal A, luminal B, HER-2 positive and triple-negative mammary cancer in female dogs.

Association between mammary tumor phenotypes and survival. Regarding the luminal A subtype, only three patients (3/42) died as a result of the disease (pulmonary metastasis). The others (39/42) were examined during the 18 months of follow-up without experiencing recurrence. For the luminal B subtype, eight dogs (8/41) died due to distant metastasis (pulmonary or hepatic). In two patients (2/41) there was a local tumor recurrence within the surgical scar. The other dogs (32/41) had no complications associated with the disease during the 18 months of follow-up. In the present study, it was observed that the majority of the patients classified as luminal A and luminal B were included in stage I of the disease and had longer survival times when compared with other phenotypes.

In the tumors with HER-2 positive, the mean survival time was 167 days. A total of seven dogs (7/10) died due to

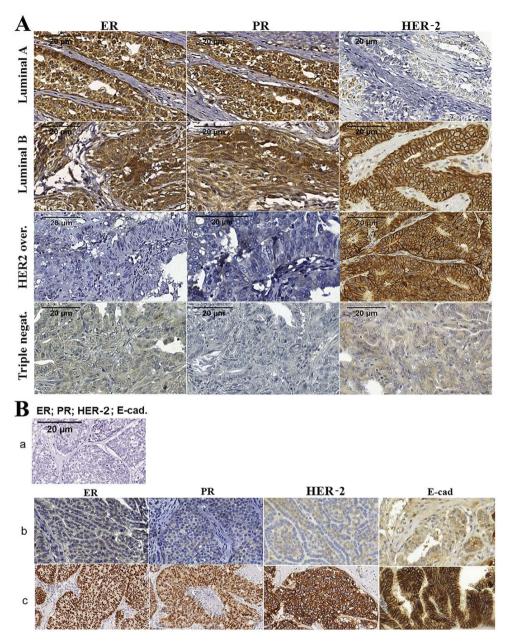


Figure 1. Immunohistochemical characteristics of mammary cancer phenotypes in bitches. (A) Photomicrograph of phenotypic profiles classified by immunohistochemical markers of ER, PR and HER-2 in canine mammary tumors. 3,3'-diaminobenzidine chromogen. Scale bar, 20  $\mu$ m. ER positive and/or PR positive and HER-2 negative signals are associated with the phenotype Luminal A. ER positive and/or PR positive and HER-2 positive signals are associated with the phenotype Luminal B. The phenotype HER-2 upregulated was characterized by ER negative, PR negative and HER-2 positive signals. ER negative, PR negative, and HER-2 negative signals are associated with the triple negative phenotype. (B) Photomicrograph of (B-a) negative human breast cancer control and (B-b) negative to low protein expression of ER, PR, HER-2 and E-cadherin in canine mammary cancer. (B-c) Human samples positive for ER, PR, HER-2 and E-cadherin. Magnification, x40. Laboratory of Molecular Investigation of Cancer (LIMC), 2019. ER, estrogen receptor; PR, progesterone receptor; HER-2, human epidermal growth factor receptor.

pulmonary metastasis, and there were no secondary complications associated with mammary cancer detected in the remaining three dogs. In the triple-negative tumor phenotype, the mean survival time associated with this profile was 274 days. A total of three dogs (3/17) did not exhibit evidence of disease complications, and 14 died from lung and/or liver metastasis. The survival curve of each phenotype is presented in Fig. 2 (P<0.0001).

Low E-cadherin expression is associated with poor prognostic outcome. Positive E-cadherin expression was observed in 80% (n=84) of mammary tumors, whereas no

staining was observed in 20% (n=21) of samples. Regarding the E-cadherin intensity, it was observed that 15 mammary tumors were scored as '+', 52 samples were scored as '++' and 17 samples were scored as '+++'.

The present study also compared the level of E-cadherin expression with the assessed phenotypic groups. A significant association was observed between the intensity of expression among the four profiles considered (P=0.03). The majority of samples with luminal A and B phenotypes presented a score of '++' and '+++'. On the other hand, tumors classified as HER-2 positive and triple-negative primarily presented scores of '0', '+' and '++' (Table III).

Table I. Antibodies and dilutions.

Antibody	Specificity	Clone	Dilution	Supplier
E-cadherin	(Mouse)	36/Ecad	1:2,000	BD Biosciences
ER	Monoclonal (mouse)	1D5	1:150	Santa Cruz Biotechnology, Inc.
PR	Monoclonal (rabbit)	SP42	1:400	Abcam
HER-2/neu	Polyclonal (rabbit)	C18	1:800	Santa Cruz Biotechnology, Inc.

ER, estrogen receptor; PR, progesterone receptor; HER-2, human epidermal growth factor receptor 2.

Table II. Distribution of frequency of clinicopathological characteristics in the mammary cancer phenotypes of female dogs.

Clinical characteristics	Luminal A, n	Luminal B, n	HER-2 positive, n	Triple-negative, n
No complications	36	32	2	4
Recurrence	1	2	0	0
Metastasis				
Cutaneous	2	0	1	0
Lung	3	5	7	12
Liver	0	2	0	1
TNM				
I	15	20	1	3
II	10	10	2	4
III	8	2	4	4
IV	5	6	1	1
V	3	1	0	4
Uninformed	1	2	2	1

TNM, Tumor-Node-Metastasis; HER-2, human epidermal growth factor receptor 2.

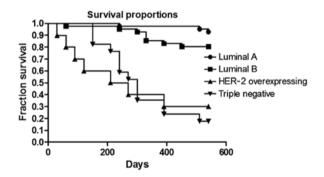


Figure 2. Kaplan-Meier survival curves representing the different phenotypes in canine mammary tumor samples. Log-rank test (P<0.001). HER-2, human epidermal growth factor receptor.

# Discussion

Previous studies have demonstrated that canine mammary cancer is a heterogeneous disease (24,26,51). Despite previous studies have tried to define histopathological classifications able to associating the type of mammary tumors and prognostic features, these attempts have failed in a number of ways; particularly in predicting disease characteristics in the early stages, in the monitoring of progress, and in the assessment of risk of recurrence and mortality in the absence of treatment (10,40,41).

The present study suggested that histopathological classification are not associated with a clear prognosis. It was observed that the same histopathological subtype may exhibit different phenotypes and, as a result, different outcomes. Therefore, it has been suggested that the anatomical and morphological evaluation of neoplastic lesions should not be considered as the only diagnostic method. In veterinary medicine, few studies investigated molecular classifications in combination with the immunohistochemical markers in canine mammary tumors (25,26,51). In addition, there are discrepancies between previous studies, which are based on a variety of markers and different techniques, making challenging to perform comparisons among previous methodologies (24-26,51,52). Notably, the main limitation of the present study was the non-standardization of the subtype and the histological grade of the neoplasms used.

Therefore, the classification proposed in the present study was based on four immunohistochemical markers (ER, PR, HER-2 and E-cadherin) that were used to define the following phenotypic groups: i) luminal A; ii) luminal B; iii) HER-2 positive; and iv) triple-negative. The use of these markers proved to be very effective in defining the phenotypic profiles of canine mammary tumors to ensure an accurate association with prognosis. This is the main result of the present study, as

**Immunostaining** Luminal A, n Luminal B, n HER-2 positive, n Triple-negative, n Negative 6 5 4 7 5 4 3 3 Weak 23 2 5 22 Moderate 9 0 1 Strong 6 9 Total 40 40 16

Table III. Distribution of frequency of intensity of E-cadherin expression in canine mammary tumors according to each phenotype.

HER-2, human epidermal growth factor receptor 2.

phenotypic classifications from previous studies use additional markers to define the panel (25,26,51).

Collectively, it was observed that the majority of canine neoplasms were classified as luminal A and luminal B, which suggested a favorable prognosis. These groups were characterized by positive expression of ERs and PRs. It has already been previously established that elevated expression levels of these receptors occurs in normal mammary glands and benign mammary tumors (3). However, mammary cancer can also exhibit expression of these receptors, which tends to have an improved prognosis while the negative staining is associated with the most malignant mammary tumors (24,52,53).

ERs are steroid receptors located in the cytoplasm and on the nuclear membrane (54,55). The activation of cytoplasmic steroidal receptors occurs via the non-genomic pathway (56). Non-genomic effects do not depend on gene transcription or protein synthesis and involve steroid-induced modulation of cytoplasmic regulatory proteins or cell membrane boundaries (57,58). Previous studies have demonstrated that the activation of this pathway influences mammary gland carcinogenesis and disease progression (59-61).

There are few studies in veterinary literature that investigated the presence and actions of ER and PR in cytoplasmic in mammary tumors (62,63). These studies revealed important information about the heterogeneity of this cancer. However, none of these focused on the prognostic value of these findings in mammary cancer. Rutteman et al (62) measured cytoplasmic ERs and PRs in normal and neoplastic mammary tissues of bitches. Rutteman et al (62) concluded that the expression of these receptors was more frequent in normal and benign neoplasm samples. Due to the results observed, they also concluded that metastases were less frequently observed in those with the same receptors. Donnay et al (63) demonstrated that poorly differentiated malignant tumors in bitches express lower concentrations of ER and PR in the cytoplasm when compared with healthy glands. Donnay et al suggested that the expression of ER and PR are greater in the most differentiated tumors and with less aggressive biological behavior. In addition, other studies (64-66) have evaluated nuclear immunoexpression of these same receptors, ER and RP, and yet the findings corroborate the results of Rutterman et al (62) and Donay et al (63).

The expression of the HER-2 protein was demonstrated to be an efficient prognostic marker (22,24,67). In the present study, there was an increased mortality rate in the luminal B group compared with the luminal A. HER-2

positive phenotype, defined by the absence of ER and PR, as well as the HER-2 expression, exhibited the worst prognosis compared with the other groups. The expression of HER-2 protein in canine mammary tumors was determined by Dutra *et al* (68) and the results are in line with the present study, as they demonstrated that upregulation of the HER-2 protein is associated with more aggressive neoplasms. However, Campos *et al* (69) observed no prognostic features associated with this marker.

In the present study, it was observed that triple-negative tumors are associated with a shorter survival time and, therefore, a worse prognosis. Kim *et al* (52) demonstrated that 18.7% of canine mammary tumors analyzed were triple-negative, and that the triple-negative phenotype was associated with a poor prognosis.

Other molecular classifications based on immunohistochemical markers for mammary tumors in female dogs were described in three previous studies. Gama et al (26) investigated an immunohistochemical panel based on five markers (ER, PR, HER-2, CK5, p63 and P-cadherin) and determined five phenotypes: Luminal A, luminal B, HER-2 positive, basal-like and null phenotype. In line with the present study, there was a significant difference between the groups regarding the prognosis and survival time, except for in the null phenotype. Sassi et al (25) proposed five immunohistochemical markers for classifying neoplasms in the luminal A, luminal B, HER-2 positive, basal-like and typical-like phenotypes. However, they only identified the luminal A subtype, luminal B and basal-like phenotypes. In contrast with the present study, there were no statistically significant differences in survival times between phenotypes (25). Im et al (51) used six antibodies (ER, HER-2, CK-14, P63, smooth muscle actin and vimentin) to define six groups: Luminal A, luminal B, HER-2 positive, basal-like and typical-like. Im et al (51) suggested that low ER expression and upregulation of HER-2 were associated with poor prognosis in canine mammary tumors.

The addition of the E-cadherin antibody to the panel ensures the classification of neoplastic cells in the epithelial profile. The association of this marker with the other groups allows the evaluation of the different behaviors among neoplasms characterized in the same phenotypic subtype. The present study demonstrated that the absence or low expression of E-cadherin was associated with more aggressive and metastatic phenotypes (triple-negative and HER-2 positive) representing an important marker that can

help to understand and better define the development and/or progression of the disease. Poorly differentiated, invasive and metastatic canine mammary carcinomas exhibited a loss of E-cadherin expression in certain tumor cell subpopulations, suggesting that altered E-cadherin expression may be an important process in malignant transformation. This hypothesis is supported by the observation that the loss of E-cadherin expression is associated with a shorter overall survival time and disease-free period (70).

Previous studies have demonstrated that the decrease in E-cadherin expression is associated with tumors of a higher invasive grade, and therefore more metastatic (13,43), in line with the results of the present study. The present study also observed that the absence or low expression of E-cadherin was found associated with more aggressive and metastatic phenotypes (triple-negative and HER-2 positive).

The proposed classification based on the phenotypic subtypes luminal A, luminal B, HER-2 positive and triple-negative was performed using immunohistochemical markers (ER, PR, HER-2 and E-cadherin), and proved to be an important prognostic tool for malignant canine mammary tumors. The classification described in the present study could be used in the field of oncology to provide valuable information about the course and evolution of mammary tumors in canine species. Notably, the main limitation of the present study was the non-standardization of the subtype and the histological grade of the neoplasms used.

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# Availability of data and materials

The datasets used during the present study are available from the corresponding author upon reasonable request.

## **Authors' contributions**

DAPDCZ conceived and designed the study. GRV performed the experiments and wrote the paper. JRL translated the main text of the article into the English language. GBG, MGM, JRL, ABDN and MTC recruited the patients and collected the samples. LBMS and RMR performed the immunohistochemistry assays. DAPDCZ revised and edited the manuscript.

## Ethics approval and consent to participate

The Ethics Committee on Animal Experimentation of the Faculty of Medicine of São José do Rio Preto

(approval no. 3231/2012) approved the present study on August 16, 2012. The Committee for Ethics in Research on Human Beings of the Faculty of Medicine of São José do Rio Preto granted the use of the human tissues.

# Patient consent for publication

Not applicable.

## **Competing interests**

The authors declare that they have no competing interests.

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