

RESEARCH ARTICLE

Evaluation of Some Systemic Inflammatory Biomarkers in Canine Malignant Mammary Tumors ^[1]

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ABSTRACT

The aim of this study is to investigate whether neutrophil-to-lymphocyte ratio (NLR), lymphocyte-to-monocyte ratio (LMR), platelet-to-lymphocyte ratio (PLR), systemic immune-inflammation index (SII), albumin-to-globulin ratio (AGR), and prognostic nutritional index (PNI) parameters could be used as biomarkers for canine malignant mammary tumors (MMTs), and the changes in these parameters according to different tumor (T), lymph node (N), and metastasis (M) stages (TNM I-II-III, TNM IV, TNM V) and the number of affected mammary glands (single, multiple). Thirty-seven with MMT and 20 healthy dogs were used in this study. Complete blood count and biochemistry analysis were performed in all dogs. Tumor material is removed by tru-cut and sent to the pathology laboratory for diagnosis. NLR, PLR, and SII values increased, and LMR and PNI values decreased in dogs with MMT. Median NLR values increased and median LMR and PNI values decreased as the TNM stage progressed. In dogs with a single MMT, median NLR, and PLR values were found to be lower than in dogs with multiple MMTs, and median LMR, SII, AGR, and PNI values were higher. The present results indicated that NLR, LMR, PLR, SII, and PNI parameters could be used as biomarkers for canine MMT. Also, NLR, LMR, PLR, SII, PNI, and AGR parameters may be valuable biomarkers that reveal the degree of systemic immune response according to different TNM stages and the number of affected mammary glands.

Keywords: Canine, Malignant mammary tumor, Inflammation, Biomarkers, Oncology

INTRODUCTION

Cancer begins when the cell becomes unable to respond to the mechanisms that control its division due to a number of structural defects in the cell ^[1]. Initially, cells that have undergone neoplastic transformation are detected by the host immune system and eliminated by various mechanisms. However, this effect of the immune system against tumors can be weak and insufficient, and even some components contribute to tumor development at the same time ^[2,3]. In fact, the tumor-associated immune response is more likely to contribute to tumor growth, progression, and immunosuppression than it is to form an effective host antitumor response ^[4]. Revealing the links between cancer and inflammation has implications

for the prevention and treatment of cancer ^[3,5]. It has been stated that cancer-related inflammation is associated with changes in circulating white blood cells and some biochemical parameters both in humans and dogs ^[6-8]. Lymphocytes have the most important role in the immune response against cancer ^[9,10]. Neutrophils and monocytes also have critical roles in antitumor immunity, they exhibit their antitumor functions directly or work with lymphocytes. However, when they gain a cancer-supportive structure, they start to exhibit quite important pro-tumor functions and contribute to tumor progression, promote metastasis, and show immunosuppressive activity ^[11,12]. Also, neutrophils can prevent the anticancer functions of lymphocytes ^[13]. Platelets become active by interacting with cancer cells and show many functions that participate



in cancer progression, metastasis, and inflammation [14]. Therefore, parameters such as neutrophil-to-lymphocyte ratio (NLR), lymphocyte-to-monocyte ratio (LMR), platelet-to-lymphocyte ratio (PLR), and systemic immune-inflammation index (SII) are calculated in many studies conducted to date. High NLR, PLR and SII values and a low LMR value are associated with advanced disease and poor prognosis. Moreover, they have been shown to be useful for the selection of the appropriate treatment method and the management of the disease in several neoplastic conditions both in humans [5,6,15-20] and dogs [21-23]. The albumin-to-globulin ratio (AGR) parameter is a useful biomarker for revealing systemic inflammation associated with malignancies and has prognostic value in various types of cancer including breast cancer in humans [24-26]. The prognostic nutritional index (PNI) is a new systemic immune-nutrition index and represents the immune and nutritional status of the host [27].

It has been demonstrated in many studies that the parameters we mentioned above can be used as biomarkers in the diagnosis, management of the disease, and prediction outcome in many cancer types, including a large population of breast cancer [15,19,24,27]. Only NLR has been reported to have a prognostic value in dogs with mammary tumors [28]. Therefore, this study aimed to investigate, firstly, whether the parameters indicating the systemic inflammatory response associated with cancer could be used as biomarkers for canine MMT, and secondly, the changes of these parameters according to different TNM stages and the number of affected mammary glands.

MATERIAL AND METHODS

Ethical Statement

The study was approved by the Ethics Committee on Animal Research of Bursa Uludag University (Approval No: 2020-03/03).

Animals

The first group of dogs in this study included 37 dogs with malignant mammary tumor (MMT) aged 3 to 16 years. The breeds of the dogs included Terrier (7), Golden Retriever (5), Crossbreed (5), Cocker (4), Doberman (2), Siberian Husky (2), Doberman Pinscher (1), Jack Russel Terrier (1), Dogo Argentino (1), Rottweiler (1), German Shepherd (1), Chihuahua (1), Pekingese (1), Beagle (1), Cane Corso (1), American Staffordshire Terrier (1), Kurzhaar (1) and Alabai (1). The second group of dogs in this study included 20 healthy dogs aged 2 to 10 years from different breeds (Crossbreed (11), Alabai (3), Golden Retriever (2), Labrador Retriever (1), Dogo Argentino (1), Border Collie (1) and Chow Chow (1)) as controls. None of the dogs had concurrent systemic inflammatory or

immune-related diseases. Two of the dogs with MMT had cancer-related cachexia. The diet of the dogs in the study was not uniform. Characteristics of dogs with MMT and the control group are summarized in *Table 1*.

Table 1. Characteristics of the dogs with malignant mammary tumors (MMTs) and the control group

Characteristics	n		Percentage (%)	
	Dogs with MMTs	Control	Dogs with MMTs	Control
Age				
≤8	17/37	17/20	45.95	85
Neutering Status				
Neutered	6/37	6/20	16.22	30
Intact	30/37	14/20	81.08	70
Remnant Ovary	1/37	0	2.70	0
Pseudopregnancy History	6/37	2/20	16.22	10
Contraception History	4/37	0	10.81	0
TNM Stage				
I-II-III	14/37		37.84	
IV	15/37		40.54	
V	8/37		21.62	
Number of Tumors				
Single	10/37		27.03	
Multiple	27/37		72.97	
Histopathological Evaluation				
Scirrhous Adenocarcinoma	9/37		24.32	
Solid Adenocarcinoma	6/37		16.22	
Tubular Adenocarcinoma	5/37		13.51	
Invasive Ductal Carcinoma	3/37		8.11	
Adenocarcinoma	3/37		8.11	
Papillary Adenocarcinoma	2/37		5.41	
Malignant Mixed Tumor	2/37		5.41	
Tubulopapillary Carcinoma	2/37		5.41	
Tubular Carcinoma	2/37		5.41	
Spindle Cell Carcinoma	1/37		2.70	
Invasive Cribriform Carcinoma	1/37		2.70	
Carcinoma	1/37		2.70	

All dogs had a general examination, complete blood count, and biochemistry analysis, in addition to that thoracic radiography and intra-abdominal ultrasonography were done in dogs with mammary tumors. Ultrasound-guided biopsy samples were obtained from mammary tumors with a 14-gauge tru-cut biopsy needle. Fine-needle biopsy was performed from local lymph nodes with a 21-gauge needle. Lidocaine Hydrochloride (Jetocaine, ADEKA, Samsun, Turkey) was used at 4 mg/kg for local anesthesia before biopsies. All biopsies were sent to the pathology laboratory. 37 dogs that were found to have MMT according to the pathological examination were included in the study. All dogs with MMT were evaluated

according to the modified TNM system [TNM I-II-III ($T_{1-2,3}N_0M_0$), TNM IV ($T_{1-2,3}N_1M_0$), TNM V ($T_{1-2,3}N_{0,1}M_1$)] as described by Goldschmidt et al.^[29]. Those with tumors in one mammary gland were grouped as “single” and those with tumors in more than one mammary gland were grouped as “multiple”.

Biochemical, Hematology Analyses and Biomarkers

Blood was taken from the vena cephalica antebrachii into EDTA tubes for complete blood count and into dry tubes for measurement of biochemical values. The blood samples centrifuged at 4000 rpm for 10 min and their serums were separated on the day of the examination. A complete blood count was performed with the “Hasvet VH5R, Automated Hematology Analyzer” (Urit, China) device. Total Protein (TP), Albumin (ALB), Alkaline Phosphatase (ALP), Glucose (GLU), Total Bilirubin (TBIL), Inorganic Phosphorus (IP), Total Cholesterol (TCHO), Gamma-Glutamyl Transferase (GGT), Alanine Aminotransferase (ALT), Calcium (Ca), Creatinine (CRE), Blood Urea Nitrogen (BUN), Globulin (GLOB), values were determined by “FUJI DRI-CHEM NX500V IC Chemistry Analyzer” (FUJIFILM, Japan) device.

The NLR was determined by dividing the neutrophil ($10^9/L$) by lymphocyte counts ($10^9/L$)^[21]. The LMR was calculated by dividing the lymphocyte ($10^9/L$) by monocyte counts ($10^9/L$)^[21]. The PLR was detected by dividing the platelet ($10^9/L$) by lymphocyte counts ($10^9/L$)^[21]. The SII was calculated by multiplying the neutrophil count ($10^9/L$) by the platelet count ($10^9/L$) and dividing by the lymphocyte count ($10^9/L$) ($N \times P/L$)^[21]. The AGR was estimated by dividing the albumin (g/dL) by globulin (g/dL)^[24]. The PNI parameter was obtained by summing 10 times serum albumin (g/dL) and 0.005 times lymphocyte count (per mm^3) ($10 \times ALB + 0.005 \times L$)^[27].

Histopathological Evaluation

Biopsy samples of mammary tumors were fixed in 10% buffered formalin and embedded in paraffin. Sections of 4 μm thickness were taken from the tissues passed through alcohol and xylol and stained with hematoxylin&eosin. Tubule and mammary alveolar formation, nuclear

polymorphism, pleomorphism, mitosis index, inflammatory infiltration, necrosis, adjacent tissue invasion, and lymph node metastasis were evaluated according to the World Health Organization criteria for canine mammary tumors^[30].

Statistical Analysis

Statistical analyses were performed using IBM SPSS Statistics 28 for Windows. Shapiro-Wilk test was used to examine whether the data were normally distributed. Nonparametric tests were used for data that did not show normal distribution. The comparison of the values of the dogs with MMT and the control group was done by using the Mann-Whitney U-test, which is one of the nonparametric tests. The Kruskal-Wallis test was used for the comparison according to the TNM stage and the number of affected mammary glands. Significance values had been adjusted by the Bonferroni correction for multiple tests. Statistical significance was set at P value <0.005.

RESULTS

The mean age of dogs with MMT and the control group was 9.59 ± 3.02 and 6.20 ± 2.31 years, respectively ($P < 0.001$). 86.48% of dogs with MMT and 45% of the control group were pure-breed dogs. The most common breeds of dogs with MMT included Terriers 18.92% (7/37), Golden Retrievers 13.51% (5/37), Crossbreeds 13.51% (5/37), and Cocker 10.81% (4/37). Characteristics of dogs with MMT are summarized in [Table 1](#). Hematological and biochemical data and reference intervals of dogs with MMT and control group are given in [Table 2](#) and [Table 3](#). Radiographic images of two dogs with lung metastases and histopathological image samples of dogs with MMTs are given in [Fig. 1](#) and [Fig. 2](#). Differences between the dogs with MMT and the control group were found statistically significant for NLR, LMR, PLR, SII and PNI parameters ($P < 0.05$). The AGR value did not show any differences between the groups ($P = 0.496$). The data are summarized in [Table 4](#).

In the comparison made according to TNM staging, the

Table 2. Hematological parameters of the dogs and the reference ranges

Variable	Dogs With CMT		Control		Reference
	Median(min:max)	Mean \pm SEM	Median(min:max)	Mean \pm SEM	
NEU $10^9/L$	8.19 (3.47:36.1)	9.36 \pm 0.90	6.05 (3.74:9.66)	6.13 \pm 0.37	2.7-9.4
MON $10^9/L$	0.61 (0.12:4.02)	0.75 \pm 0.11	0.32 (0.09:0.64)	0.34 \pm 0.03	0.1-1.3
LYM $10^9/L$	1.55 (0.78:4.47)	1.79 \pm 0.15	2.48 (1.15:4.91)	2.53 \pm 0.19	0.9-4.7
PLT $10^9/L$	349 (147:693)	377.78 \pm 24.51	263 (172:384)	265.95 \pm 12.35	186-545

Data expressed as median (min:max) and mean \pm SEM. NEU=neutrophils, MON=monocytes, LYM=lymphocytes, PLT=platelets.

Table 3. Biochemical parameters of the dogs and the reference ranges					
Variable	Dogs With MMTs		Control		Reference Values
	Median(min:max)	Mean±SEM	Median(min:max)	Mean±SEM	
TP g/dL	6.8 (5.6:8.2)	6.71±0.10	6.5 (5.9:7.1)	6.48±0.08	5.5-7.2
ALB g/dL	3.4 (2.3:4.5)	3.39±0.08	3.4 (3:4)	3.39±0.06	3.2-4.1
GLOB g/dL	3.1 (2.5:5.4)	3.29±0.11	3.1 (2.6:3.7)	3.09±0.06	1.9-3.7
ALP U/L	56 (14:287)	80.31±11.35	38 (19:86)	43.63±4.93	7-115
GLU mg/dL	104 (64:134)	104.2±2.38	107.5 (70:123)	101.95±3.50	68-104
TBIL mg/dL	0.3 (0.2:0.6)	0.28±0.02	0,2 (0.2:0.4)	0.24±0.02	0-0.2
IP mg/dL	3.6 (1.5:6.5)	3.61±0.17	3.5 (2.9:4.6)	3.65±0.10	2.7-5.4
TCHO mg/dL	287 (169:437)	302.18±15.49	165 (108:275)	175±12.59	136-392
GGT U/L	<10	<10±0	<10	<10±0	0-8
ALT U/L	41.5 (10:149)	45.16±3.94	36.5 (20:75)	41.25±3.25	17-95
CA mg/dL	10.6 (9.3:12.1)	10.65±0.11	11.1 (10.2:11.9)	11.15±0.11	9.4-11.1
CRE mg/dL	0.75 (0.41:1.6)	0.81±0.05	0.69 (0.43:0.97)	0.66±0.031	0.6-1.4
BUN mg/dL	14.05 (5:29)	14.62±1.09	13 (5:19.9)	12.43±1.01	9-26

Data expressed as median (min:max) and mean ± SEM. ALB=albumin, GLOB=globulin, APL=alkaline phosphatase, GLU=glucose, TBIL=total bilirubin, IP=inorganic phosphorus, TCHO=total cholesterol, GGT=gamma-glutamyl transferase, ALT=alanine aminotransferase, CA=calcium, CRE=creatinine, BUN=blood urea nitrogen

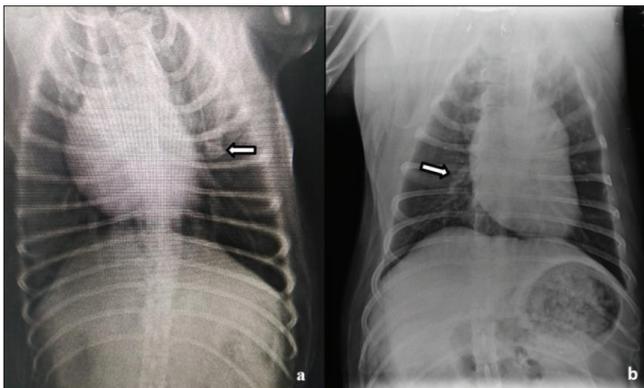


Fig 1. Ventrodorsal radiographic views of the thorax in a 5 years old Terrier (a) and in an 8 years old Cane Corso (b) with MMTs

NLR value did not show statistical significance between the control group and TNM I-II-III stage ($P=0.069$), while the differences between each of TNM IV and TNM V and the control group were significant ($P<0.001$, $P=0.012$, respectively). The LMR values were found to be statistically significantly lower in each of the TNM I-II-III, TNM IV, and TNM V stages compared to the control group ($P=0.032$, $P<0.001$, $P=0.006$, respectively). The PLR values were determined significantly higher in TNM I-II-III and TNM IV stages versus the control group ($P=0.03$, $P=0.001$, respectively). However, the difference between the PLR values of the TNM V group and the control group was not significant ($P=0.076$). The SII values were significantly higher in TNM I-II-III, TNM IV, and TNM V stages versus the control group ($P=0.018$, $P<0.001$, $P=0.038$, respectively). The overall test did not

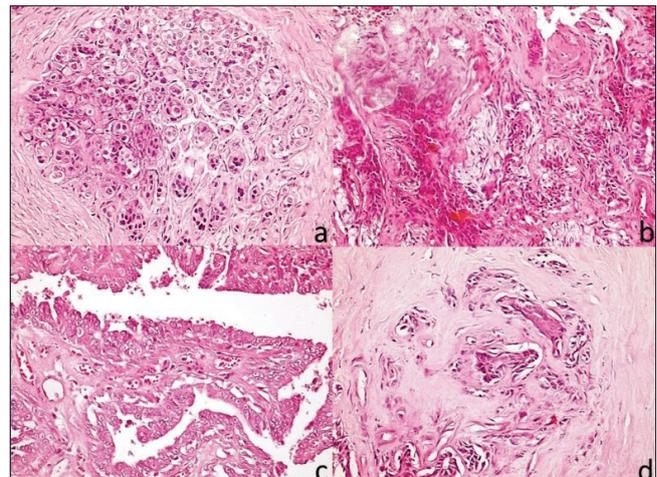


Fig 2. a- Solid Adenocarcinoma. Neoplastic cells exhibit a pronounced pleomorphism and fill the alveolar lumen, b- Malignant Mixed Tumor. The neoplasm contains multiple clusters of carcinoma cells as well as regions characterized by the proliferation of myoepithelial cells, c- Tubulopapillary carcinoma. Neoplastic cells have a vesicular appearance and are binucleated. Tubular epithelia have formed multiple layers, d- Scirrhous Adenocarcinoma. Excessive increase in fibrous stroma led to deterioration of lobular structure. Alveoli and ducts are not visible

show significant differences between the groups for the AGR parameter ($P=0.344$). The PNI values did not show statistically significant differences between TNM I-II-III and TNM V stages versus the control group ($P>0.05$), but the difference in the PNI value between TNM IV and control group was significant ($P=0.039$). There was no significant difference in the biomarkers assessed among TNM I-II-III, TNM IV, and TNM V stages (Table 5).

The NLR, LMR, PLR, and SII values were found to be

Table 4. Comparison of dogs with malignant mammary tumors (MMT) and control group

Parameters	Dogs with MMTs		Control		P Value
	Median (min : max)	Mean ± SEM	Median (min : max)	Mean ± SEM	
NLR	5.01 (1.56:29.35)	6.34±0.80	2.77 (1.12:4.30)	2.63±0.20	<0.001
LMR	2.68 (0.31:12.83)	3.89±0.56	6.99 (3.82:27.56)	9.26±1.44	<0.001
PLR	242.11 (47.09:689.66)	262.79±25.92	107.28 (48.73:236.36)	117.80±10.66	<0.001
SII	1791.98 (383.58:12855.12)	2574.62±420.64	724.18 (199.87:1342.55)	713.05±70.23	<0.001
AGR	1.07 (0.52:1.73)	1.08±0.05	1.06 (0.92:1.54)	1.11±0.03	0.496
PNI	41.93 (31.9:56.65)	43.15±6.59	45.38 (39.75:58.55)	46.47±1.08	0.022

Data expressed as median (min:max) and mean ± SEM. NLR=neutrophil-to-lymphocyte ratio, LMR=lymphocyte-to-monocyte ratio, PLR=platelet-to-lymphocyte ratio, SII=systemic immune-inflammation index, AGR=albumin-to-globulin ratio, PNI=prognostic nutritional index.

Table 5. Comparisons according to TNM stages of the dogs with malignant mammary tumors.

Parameters	TNM I-II-III		TNM IV		TNM V		Control		P Value
	Median (min : max)	Mean ± SEM	Median (min : max)	Mean ± SEM	Median (min : max)	Mean ± SEM	Median (min : max)	Mean ± SEM	
NLR	4.66 ^{ab} (1.86:14.55)	4.84 ±3.21	5.83 ^a (3.02:29.35)	7.97±1.70	6.51 ^a (1.56:8.62)	5.9±0.92	2.77 ^b (1.12:4.30)	2.63±0.20	<0.001
LMR	3.07 ^a (0.81:12.83)	5.00±1.08	2.54 ^a (0.31:10.59)	2.99±0.60	2.27 ^a (1.07:12.25)	3.63±1.34	6.99 ^b (3.82:27.56)	9.26±1.44	<0.001
PLR	200.37 ^a (47.76:689.66)	252.21±48.37	255.45 ^a (47.09:617.50)	286.08±40.52	254.74 ^{ab} (55.03:431.03)	237.64±44.23	107.28 ^b (48.73:236.36)	117.80±10.66	0.001
SII	1478.84 ^a (525:8731.03)	2159.07±601.15	2165.96 ^a (538.24:12855.12)	3272.57±840.97	2082.66 ^a (383.58:3723.86)	1993.18±399.91	724.18 ^b (199.87:1342.55)	713.05±70.23	<0.001
AGR	1.19 (0.76:1.73)	1.74±0.08	0.94 (0.52:1.54)	1.02±0.10	0.94 (0.84:1.25)	1.01±0.07	1.06 (0.92:1.54)	1.11±0.03	0.344
PNI	43.88 ^{ac} (37:56.65)	45.56±1.61	41.25 ^{bc} (31.9:54.90)	41.10±1.87	40.80 ^{bc} (33.95:54.35)	42.10±2.54	45.38 ^a (39.75 8.55)	46.47±1.08	0.029

^{a-c} Different superscripts indicate values that within the row are significantly different. Data expressed as median (min:max) and mean ± SEM. NLR=neutrophil-to-lymphocyte ratio, LMR=lymphocyte-to-monocyte ratio, PLR=platelet-to-lymphocyte ratio, SII=systemic immune-inflammation index, AGR=albumin-to-globulin ratio, PNI=prognostic nutritional index

Table 6. Comparisons according to the number of affected mammary glands.

Parameters	Single		Multiple		Control		P Value
	Median (min : max)	Mean ± SEM	Median (min : max)	Mean ± SEM	Median (min : max)	Mean ± SEM	
NLR	4.73 ^a (2.17:8.62)	4.88±0.72	5.34 ^a (1.56:29.35)	6.87±1.06	2.77 ^b (1.12:4.30)	2.63±0.20	<0.001
LMR	3.23 ^a (1.07:11.00)	4.70±1.19	2.68 ^a (0.31:12.83)	3.59±0.63	6.99 ^b (3.82:27.56)	9.26±1.44	<0.001
PLR	232.58 ^a (97.40:382.58)	240.23±32.57	242.11 ^a (47.09:689.66)	271.15±33.60	107.28 ^b (48.73:236.36)	117.80±10.66	<0.001
SII	1889.92 ^a (525.00:3723.86)	1898.28±314.00	1734.58 ^a (383.58:12855.12)	2825.12±560.49	724.18 ^b (199.87:1342.55)	713.05±70.23	<0.001
AGR	1.12 (0.52:1.62)	1.09±0.12	1.03 (0.61:1.73)	1.08±0.06	1.06 (0.92:1.54)	1.11±0.03	0.781
PNI	43.10 (33.95:56.30)	43.66±2.26	41.70 (31.90:56.65)	42.96±1.33	45.38 (39.75:58.55)	46.47±1.08	0.066

Different superscripts indicate values that within the row are significantly different. Data expressed as median (min:max) and mean ± SEM. NLR=neutrophil-to-lymphocyte ratio, LMR=lymphocyte-to-monocyte ratio, PLR=platelet-to-lymphocyte ratio, SII=systemic immune-inflammation index, AGR=albumin-to-globulin ratio, PNI=prognostic nutritional index

significantly different between dogs with a single MMT and control group ($P=0.037$, $P=0.024$, $P=0.009$, and $P=0.002$, respectively). Similarly to that, the same values of the dogs with multiple MMTs and the control group were found significantly different ($P<0.001$, $P<0.001$, $P=0.001$, and $P<0.001$, respectively). In dogs with a single MMT, the median NLR and PLR values were lower and the median LMR and SII values were higher than in dogs with multiple MMTs, but no statistical significance was found ($P>0.05$). The overall test did not show significant differences between the groups for the AGR and PNI parameters ($P=0.781$, $P=0.066$, respectively) (Table 6).

DISCUSSION

There are strong similarities between human breast cancer (HBC) and canine mammary tumors [31-33]. Also, neoadjuvant chemotherapy can be used in the treatment of canine mammary tumors similar to that in humans [34]. As in HBC [6,11,35], it has been noted that there are some remarkable changes in blood values in the presence of mammary tumors in dogs [7,8]. In veterinary medicine, it has been stated that NLR, LMR, PLR, and AGR, parameters may be potential biomarkers in certain malignancies of dogs [21-23,28]. But until recently, no literature data were evaluating LMR, PLR, SII and PNI parameters in dogs with MMTs. Uribe-Querol et al. [28] reported that a high NLR value ($NLR<5$) before treatment was associated with a lower survival rate in dogs with mammary tumors. They also stated that NLR could be used as a prognostic marker for disease severity, but AGR value did not show any predictive value on tumor malignancy. In this study, we found that in dogs with MMT, NLR, PLR, and SII values were high, and LMR and PNI values were low as in human breast cancer [15,18,27]. However, in our dogs with MMT, there was no significant difference compared to our control group in the AGR parameter [24-26]. In the study by Lallo et al. [36], it was stated that AGR values were lower in malignant MMTs. Unlike this, the AGR value did not differ in dogs with MMT when compared with healthy ones in the study by Uribe-Querol et al. [28] and in this study. We recommend evaluating the AGR parameter in larger populations.

Median NLR, PLR, LMR, and SII values show differences among TNM stages in HBC. It has been said that NLR, PLR, LMR, and SII parameters can be valuable and guide in the staging of HBC [18]. Low AGR has been found to be associated with advanced-stage of HBC and low PNI parameter has been found to be associated with advanced disease [24,27]. High NLR values have been associated with advanced or aggressive HBC [5,15,20]. It has been shown that NLR values increase as the disease progresses in TNM stages of HBC [17,18]. In our study, although not statistically significant, the median NLR values increased as TNM

stages progressed (4.66, 5.83, 6.51, respectively). Low LMR is correlated with advanced disease and TNM stages in HBC [18,20]. Compatible with this, median LMR values decreased as TNM stages progressed (3.07, 2.54, 2.27, respectively) in our study, but this decrease was not statistically significant. It was shown that the PLR parameter was correlated with advanced disease and TNM stages [18,20], and another study reported that it was associated with lymph node metastasis but not with advanced T stages [16]. Unlike them, Elyasinia et al. [17] reported that there was no relationship between PLR values and different TNM stages of HBC. Similarly, PLR values did not show a significant difference among TNM stages in the dogs with MMT. The SII parameter has been found to be associated with the advanced TNM stage in HBC [18,19]. The median SII values in TNM IV and V stages were found to be higher than the median SII value in the TNM I-II-III stages in our study. Low AGR has been found to be associated with the advanced stage of HBC [24]. In our study, the highest median AGR value was observed in the TNM stage I-II-III group, in which no metastases have formed. A low PNI parameter has been found to be associated with advanced disease in HBC [27]. Although there was no statistically significant difference in our study, median PNI values decreased as TNM stages progressed (43.88, 41.25, 40.8, respectively). Similar changes were detected in NLR, LMR, PLR, and PNI parameters at TNM stages in dogs with MMT as in HBC. These biomarkers reveal the balance between the immune system and systemic inflammation. In the early stages of the disease, tumors can be detected and destroyed by the immune system. As the disease progresses, the immunogenic capacity of the tumor decreases and its inflammatory capacity increases. As detectable tumors develop, cancer cells develop different mechanisms that mimic peripheral immune tolerance to avoid tumorocidal attack [2,37]. In our study, although the changes in SII and AGR parameters according to TNM stages showed some differences from HBC, median values in TNM I-II-III stages suggested that less systemic inflammatory response occurred in the early stages of canine MMT (Table 5).

Lymph node involvement seems to be the most important factor in predicting prognosis using systemic inflammatory parameters in HBC [16,19,20,25,27]. In our study, it was found that there was no significant increase in NLR values up to TNM stage IV, where the disease affected the lymph nodes for the first time. The highest NLR, SII, and median PLR, SII values, with the lowest LMR, AGR, and PNI values were obtained in TNM stage IV (Table 5). The most significant P values were obtained for the NLR, LMR, PLR, and SII parameters compared to the control group and TNM stage IV. Interestingly, the PNI parameter showed a significant difference only when TNM stage IV

and the control group were compared (Table 5). However, no statistical difference was found between TNM IV and V groups. According to the current literature, the rate of bilateral HBC development humans is low [38] but, multiple tumors are more common in dogs [39,40]. In dogs with a single MMT, median NLR, and PLR values were found to be lower than in dogs with multiple MMTs, and median LMR, SII, AGR, and PNI values were higher in our study (Table 6). These results suggest that the systemic inflammatory response may be higher when the disease metastasizes to other mammary glands than when it is localized in a single mammary gland. More research should be done about this subject in larger populations.

Advanced clinical staging is known to be associated with hematologic parameters and provide prognostic information for canine mammary tumors [8]. The data of this study showed that some combinations (NLR, LMR, PLR, SII, and PNI) of hematological and biochemical data routinely measured for systemic inflammatory response, varied between healthy dogs and dogs with malignant mammary tumors. It is also suggested that these biomarkers could be used as biomarkers in different TNM stages of MMTs and in cases where the disease metastasizes from the mammary gland of the primary tumor to other mammary glands. However, further studies are needed to determine the value of these biomarkers in determining the prognosis of the disease, choice of treatment modality, or prediction of response to chemotherapy.

Availability of Data and Materials

Data supporting these findings are available upon request from the corresponding author (D. NAK) on reasonable request.

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Conflict of Interest

The authors declare no conflict of interest.

Ethical Statement

The study was approved by the Ethics Committee on Animal Research of Bursa Uludag University (Approval No: 2020-03/03).

Author Contributions

D.N. and Z.M.E. wrote the manuscript, conducted experiments and interpreted the results. M.O.O. and Z.A.K performed the histopathological examination. F.E.K. and T.A. made a significant contribution to and supported the experiments. D.K. and O.G. participated in the experiments. Y.N. and all authors critically reviewed and revised the manuscript draft and approved the final version for publication.

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