

Tissue fibrosis and its correlation with malignancy in canine mammary tumors^a

Fibrosis tisular y su correlación con malignidad en neoplasias mamarias caninas

Fibrose tecidual e sua correlação com a malignidade em neoplasias mamárias em cadelas

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Abstract

Background: Fibrosis is present in several pathologies associated with mammary carcinogenesis. **Objective**: To evaluate and quantify the fibrosis present in malignant and benign mammary neoplasms in bitches. **Methods**: Eighty-three samples were divided according to histopathological diagnosis into benign (n=21) and malignant (n=62) neoplasms. Haematoxylin-eosin and Masson's trichrome were used to locate the connective tissue, and the extent of fibrosis was assessed with image software. **Results**: Benign neoplasms were classified into adenomas (cystic, complex, and tubular), benign mixed tumor, and ductal and lobular hyperplasia. Malignant neoplasms were classified as carcinomas (complex, mixed tumor, *in situ* tubular, tubulopapillary, and solid). Grade I was the most prevalent histopathological class, followed by grade II and III. Fibrosis was classified as severe, moderate, or discrete. No significant (p>0.05) difference (p=0.028) was found for fibrosis percentage between histopathological subtypes of tumors. The benign subtype of lobular hyperplasia presented differences between cystic adenoma and benign mixed tumor. The *in situ* malignant tubular carcinoma subtype presented differences between solid and tubulopapillary carcinoma. **Conclusions:** Fibrosis in canine mammary tumors can be estimated with Massons's trichrome staining.

Keywords: biological biomarkers, breast cancer, dogs, histological stain, histopathology, Masson's trichrome.

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Resumen

Antecedentes: La fibrosis está presente en diversas patologías asociadas con carcinogénesis mamaria. **Objetivo:** Analizar si existe una correlación entre fibrosis tisular y malignidad en tumores mamarios caninos. **Métodos:** 83 muestras de tejido mamario fueron divididas en masas benignas (n=21) y malignas (n=62), de acuerdo con sus características histopatológicas. En estas muestras se utilizaron las coloraciones de hematoxilina-eosina y tricromo de Masson para localizar el tejido conectivo y se analizó la proporción y cuantificación de fibrosis en los mismos con un software de imagen especializado. **Resultados:** Las masas mamarias benignas se diagnosticaron como adenomas (quístico, complejo y tubular), tumor mixto benigno, e hiperplasia ductal y lobular; las masas malignas, como carcinomas (complejo, tumor mixto, tubular *in situ*, túbulopapilar y solido). Los tumores grado I fueron los más prevalentes, seguidos por los grados II y III. La fibrosis se clasifico como discreta, moderada o severa. No se observó diferencia (p>0,05) en el porcentaje de fibrosis entre neoplasias benignas y malignas. No obstante, el porcentaje de fibrosis mostro diferencias (p=0,028) entre subtipos tumorales. La hiperplasia lobular fue diferente en relación a los adenomas quísticos y tumor mixto mamario. El subtipo maligno carcinoma tubular *in situ* fue diferente respecto a los carcinomas complejo, mixto maligno, sólido, y tubulopapilar. **Conclusiones:** La fibrosis en los tumores mamarios caninos se puede estimar por la tinción con tricromo de Masson.

Palabras claves: biomarcadores tumorales, cancer de mama, caninos, coloración histológica, histopatología, Masson's trichrome.

Resumo

Antecedentes: A fibrose participa em diversas patologias e ainda possui função adicional associada à carcinogênese mamária. **Objetivo:** Objetivou-se avaliar a fibrose e correlacionar com a malignidade nas neoplasias mamárias em cadelas. **Métodos:** 83 amostras foram divididas em dois grupos baseado no diagnóstico histopatológico: neoplasias benignas (n=21) e neoplasias malignas (n=62). Hematoxilina e eosina e Tricômico de Masson foram usadas para vizibilização de tecido conjuntivo e avaliação de fibrose através de programa de imagens. **Resultados:** As neoplasias benignas foram: adenoma (cístico, complexo e tubular), tumor misto benigno, hiperplasia ductal e lobular. As malignas foram: carcinoma complexo, em tumor misto maligno, tubular *in situ*, tubulopapilar e sólido. O grau histopatológico prevalente foi grau I, seguido do grau II e III. A fibrose nas neoplasias mamárias malignas foi classificada como severa, moderada e discreta. Não houve diferença significativa (p>0,05) na porcentagem de fibrose em relação ao diagnóstico histopatológico. O subtipo benigno hiperplasia lobular apresentou diferença entre o adenoma cístico e o tumor misto benigno. O subtipo maligno carcinoma tubular *in situ* apresentou diferença entre carcinoma complexo, misto maligno, sólido e tubulopapilar. **Conclusões:** A fibrose nos tumores mamários caninos pode ser estimada através da coloração de tricrômio de Masson.

Palavras-chaves: biomarcadores tumorais, cães, câncer de mama, coloração histológica, histopatologia, Tricômico de Masson.

Introduction

Fibrosis is an active biosynthetic process characterized by unregulated accumulation of extracellular matrix (ECM) in chronic ischemic lesions as a result of chemical or toxic agents, viral and non-viral infections, physical damage, and immunological reactions. Tumor stiffness is a feature associated with the tumoral microenvironment and recent studies have indicated that it may influence neoplastic progression involving ECM (Thannickal, 2004; Wynn, 2008; Guarino *et al.*, 2009). Tissue stiffness is clinically considered an oncogenic risk factor when correlated to other conditions, as seen in dense mammary tissue and liver cirrhosis. In humans, tumor stiffness is one of the clinical features examined during palpation of the mammary gland (Liu *et al.*, 2016) as the neoplastic tissue is stiffer than normal breast tissue and the stiffening process begins at the early stages of tumoral development (Barr *et al.*, 2015). Chronic fibrosis and inflammation has been associated with mammary tumors, indicating poor prognosis for the patients (Kalliuri and Zeisberg, 2006; Carvalho *et* *al.*, 2016). Studies have demonstrated that as fibrosis progresses, the signaling of pre-neoplastic cells with pro-inflammatory cytokines gradually increases (Kalliuri and Zeisberg, 2006).

Mammary tumors are the most common neoplasms in entire bitches and approximately 40 to 50% are considered malignant (Soremo, 2003; Cassalli *et al*, 2014; Visan *et al.*, 2016). Thus, canine mammary tumors can be used as a relevant animal model for breast cancer studies in women (Klopfleisch *et al.*, 2011; Queiroga *et al.*, 2011; Feliciano *et al*, 2012; Jensen-Jarolim *et al.* 2015; Visan *et al.*, 2016; Carvalho *et al.*, 2016). In spite of this, the correlation between mammary tumors and the degree of fibrosis in bitches has yet to be evaluated.

Therefore, it is hypothesized that canine malignant mammary tumors with high histopathological grade and short evolution period have more fibrosis than benign tumors. Thus, the aim of this study was to evaluate and quantify fibrosis present in malignant and benign mammary neoplasms in bitches.

Materials and methods

Ethical considerations

The methods used were in agreement with the Ethics Committee on the Use of Animals (CEUA, process N. 023705/12) of UNESP.

Type of study, location, animals and samples

A retrospective study was conducted using mammary tumor samples from animals admitted to the Serviço de Obstetrícia Veterinária da Faculdade de Ciências Agrárias e Veterinárias, Universidade Estadual Paulista (UNESP), Jaboticabal Campus, Brazil, from January 2013 to February 2014.

Eighty-three samples of mammary tumors from bitches of different ages and breeds were used in this study. One sample was selected per animal. In animals that presented two or more histological types, only the sample from the most histologically aggressive type was selected.

At the time of patient admittance, the tumors were examined and tumoral nodule diameter measured using a pachymeter (Lee Tools, Santo André, SP, Brazil). Nodule diameter was classified (T) according to the World Health Organization (Owen, 1980) into T1 (\leq 3 cm), T2 (between 3 and 5 cm), and T3 (\geq 5 cm). Nodule size, animal age, and evolution time of tumor were tested for correlation with degree of tumoral fibrosis. Following mastectomy, fragments of mammary tumors were collected and processed for histopathological diagnosis under light microscopy. Samples were fixed in 10% phosphate buffered formalin (pH 7.4) for 24 hours, dehydrated in decreasing alcohol solutions, diaphonized in xylene, and embedded in paraffin. Sections (5 μm) were stained with haematoxylin-eosin (HE) for further identification of main morphological alterations.

Neoplasms were classified under light microscopy, according to criteria by Misdorp *et al.* (1999). Histopathological grading was performed according to the methodology by Elston and Ellis (1998), and the samples distributed into two experimental groups: Group 1 (G1, benign neoplasms, n=21) and Group 2 (G2, malignant neoplasms, n=62).

Histopathological sections stained with Masson's trichrome (MT) were used to view connective tissue and determine fibrosis degree. A score was generated according to the extent of fibrosis in the samples. Fibrosis degree was determined by the Banff Classification (Solez *et al.*, 1993), as follows: Grade 1 (discrete, 6 - 25% of the sample), Grade 2 (moderate, 25.01 - 50% of the sample), and Grade 3 (severe, \geq 50.01% of the sample).

To evaluate the extent of fibrosis, five fields from each slide were randomly photographed (20x objective) and analyzed using the Image-Pro Plus16, version 4.5 (Medica Cybernectics, Inc., Rockville, MD, USA) processing software, which automatically calculates the percentage of staining in the areas marked in the field. The results for each slide were obtained by calculating the mean percent of the areas marked in the five fields (Figure 1).

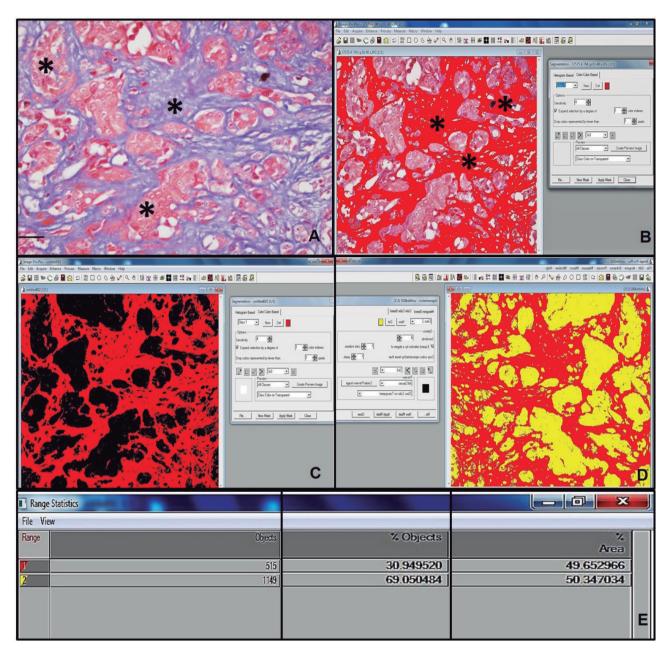


Figure 1. Histopathological photomicrograph of the mammary gland in a bitch. A: Solid carcinoma (*), Masson's trichrome (MT), bar=100 μ m. B: Histopathological photograph preparation of the mammary gland of a bitch through Image-Pro Plus16 software. Fibrosis area marked by Image-Pro Plus 16 image software (*). C: Marked fibrosis area analysis (red), chosen automatically by the image program. D: Calculation of marked fibrosis area (red) and background color (yellow). E: Result of automatic calculation of fibrosis area in solid carcinoma.

Statistical analysis

The percentage of fibrosis was compared between benign and malignant neoplasms regarding histopathological diagnosis, age of the animal, tumor evolution time, and histopathological grade using the non-parametric Kruskal-Wallis test. The correlation between histopathological grade, age, tumor evolution time, and malignancy was performed by Pearson's test. Once the median of the fibrosis percentage in malignant neoplasms was calculated, the sensitivity, specificity, positive predictive value (PPV), negative predictive value (NPV), and accuracy of the diagnostic test for malignancy were determined. The relationship between fibrosis percentage and probability of the diagnosis being malignant was also tested by binary logistic regression. Differences were considered significant at p<0.05.

Results

Out of the eighty-three samples used in this study, 21 (25.30%) were classified as benign and 62 (74.70%) as malignant neoplasms. The benign tumors were further classified into cystic adenomas (1, 1.20%), complex adenoma (4, 4.81%), tubular adenoma (2, 2.40%), benign mixed tumor (4, 4.81%), ductal hyperplasia (4, 4.81%), and lobular hyperplasia (6, 7.22%). The malignant neoplasms were classified into complex carcinoma (7, 8.43%), malignant mixed tumor (21, 25.30%), *in situ* tubular carcinoma (3, 3.61%), tubulopapillary carcinoma (18, 21.68%), and solid carcinoma (13, 15.66%) (Figure 2). Grade I was the most prevalent histopathological grade in malignant neoplasms (27, 43.55%), followed by grade II (25, 40.32%) and III (10, 16.13%).

Image analysis of the fibrotic tissue area in 45 samples revealed prevalence of severe fibrosis in malignant mammary neoplasms, followed by a lower occurrence of moderate and discrete fibrosis. Moderate fibrosis was also prevalent in benign mammary tumors, followed by severe and discrete fibrosis (Table 1). Fibrosis degree was classified as discrete, moderate or severe. When evaluating fibrosis intensity in canine breast neoplasms (discrete, moderate or severe) between the experimental groups, no statistical difference was found (p=0.174). Mean value (\pm SD) in group 1 was 3.0 \pm 1.0, and group 2 was 2.5 \pm 1.0.

Table 1. Frequency and percentage (%) of fibrosis degrees in benign and malignant mammary tumors in bitches.

Malignant Degree of Fibrosis	Frequency	%	Benign Degree of Fibrosis	Frequency	%
Discrete	05	6.02	Discrete	01	1.20
Moderate	12	14.46	Moderate	12	14.46
Severe	45	54.22	Severe	08	9.64
Total	62	74.70		21	25.30

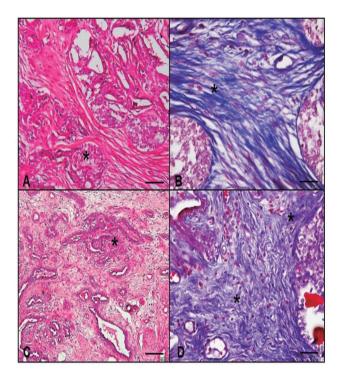


Figure 2. Histopathological photomicrograph of the mammary gland in a bitch. A: Malignant mixed tumor (*), HE, bar 100 μ m. B: Area of fibrosis in malignant mixed tumor (*), MT, 100 μ m bar. C: Complex carcinoma (*), HE, bar 100 μ m. D: Area of fibrosis in complex carcinoma (*), MT, bar 100 μ m. MT: Masson's trichrome, HE: haematoxylin-eosin.

No significant (p=0.134) difference in the percentage of fibrosis between malignant and benign tumors was observed. Mean (\pm SD) percentage of fibrosis in malignant and benign neoplasms was 58.2 \pm 24.50 and 49.9 \pm 19.10, respectively.

Fibrosis percentage was positively correlated with histopathological grade of tumors (p=0.035, r=0.23), which in turn had a positive correlation with age (p=0.003, r=0.32). In malignant tumors, severe fibrosis (\geq 50.01% of the sample) was present, in decreasing order, in samples of *in situ*, tubular carcinoma, malignant mixed tumor, solid carcinoma, complex carcinoma, and tubulopapillary carcinoma. Solid carcinoma, complex carcinoma, and tubulopapillary carcinoma presented discrete fibrosis (\leq 25% of the sample). The benign histological types with severe fibrosis were mixed tumor, cystic adenoma, and ductal hyperplasia. Tubular adenoma, complex adenoma, and lobular hyperplasia presented moderate fibrosis (25.01 - 50% of the sample) and complex adenoma showed discrete fibrosis ($\leq 25\%$ of the sample).

Fibrosis percentage was different (p=0.0284) between histopathological subtypes, as shown in Table 2.

Table 2. Median \pm interquartile range (IQR) of tissue fibrosis (%) in the different histopathological subtypes of benign and malignant canine mammary tumors.

Groups	Median ± IQR		
Group I – Benign			
Cystic Adenoma	56.30 ± 0^{ab}		
Complex Adenoma	56.55 ± 0.00^{ab}		
Tubular Adenoma	42.45 ± 2.09^{b}		
Ductal Hyperplasia	57.75 ± 6.95 ^{ab}		
Lobular Hyperplasia	39.35 ± 13.10 ^a		
Benign Mixed Tumor	63.60 ± 4.30^{ab}		
Group II – Malignant			
Complex Carcinoma	47.70 ± 19.15 ^{ab}		
Malignant Mixed Tumor	63.45 ± 15.00 ^{ab}		
<i>In situ</i> Tubular Carcinoma	73.30 ± 3.90^{a}		
Solid Carcinoma	61.75 ± 10.45 ^{ab}		
Tubulopapillary Carcinoma	51.10 ± 16.80 ^a		

Different superscript letters (^{a, b}) within the same column indicate significant difference between histological types (p<0.05) by the Dunns' test. IQR: \pm interquartile range.

Regarding tumoral evolution time, the nodules appeared between one and 36 months. Mean evolution time in benign and malignant neoplasms was 7.76 and 7.06 months, respectively. There was no significant difference in the percentage of fibrosis between different evolution times (p=0.14). Furthermore, fibrosis percentage was negatively correlated with tumoral evolution time (p=0.008, r=-0.3), suggesting that evolution time does not determine the chronicity of the process.

As the median percentage of fibrosis in malignant neoplasms was 58%, the sensitivity and specificity of the test to estimate tumor malignancy were evaluated. The test showed 75% prevalence, 45% sensitivity, 86% specificity, 90% positive predictive value (PPV), and 34% negative predictive value (NPV). The positive and negative verisimilitude were 94 and 24%, respectively. Test accuracy was 38%.

There was no significant (p=0.528) difference in the percentage of fibrosis between the different nodule

sizes, with mean (\pm SD) percentage of 54.93 \pm 21.68, 57.51 \pm 22.69, and 60.36 \pm 31.63 for T1, T2, and T3 mases, respectively. There was no positive correlation between tumoral size and fibrosis percentage, animal age, histopathological grade, or tumoral evolution time, when they were compared separately.

Binary logistic regression was used to test the relationship between fibrosis percentage and probability of a malignant diagnosis. Fibrosis percentage was predictive of malignancy in only 46% of the tumor samples (p=0.074).

Discussion

Histopathological analysis in the present study revealed that malignant neoplasms corresponded to 74.70% of the samples, in agreement with Oliveira et al (2003). Our findings also agreed with Oliveira Filho et al (2010), Sorenmo et al (2009), Feliciano et al (2012), and Soler et al (2016), who reported similar findings (73.4, 79.9, 60, and 65.38%; respectively). The frequency of benign and malignant neoplasms reported in the literature varies markedly due to different classification methods of mammary tumors and lack of standard criteria to differentiate tumor types (Cassali et al., 2003). In other countries, the percentages of malignant neoplasms reported are often lower than 54% (Itoh et al., 2005; Sorenmo et al., 2009; Lana et al., 2013; Goldshimit et al., 2017). On the other hand, some studies in Brazil have reported the prevalence of malignant neoplasms ranging from 60 to 91% (Nardi et al., 2002; Oliveira et al., 2003; Feliciano et al., 2012). This difference is probably due to the high prevalence of malignant neoplasms in Brazilian studies and the extensive time elapsed from tumor appearance to clinical evaluation (Oliveira et al., 2003). Furthermore, there is evidence that prolonged evolution time enables progression of benign into malignant tumors (Sorenmo et al., 2009).

A high prevalence of malignant mixed tumor and tubulopapillary carcinoma amongst malignant mammary neoplasms was observed, in agreement with Cintra *et al.* (2014) who reported prevalence of malignant mixed tumor when using Moulton's (1990) classification. However, Soler *et al.* (2016), using the classification by Goldshimidt *et al.* (2011), reported higher incidence of complex and simple carcinoma in mammary tumors. In the classification by Cassali *et al.* (2011), there were some differences to the classification of malignant mixed tumors. This histological subtype (malignant mixed tumor) is separated from the mixed tumor carcinoma subtype due to its reduced invasiveness and better prognosis.

According to McPhail and Robinson (2010), presence of fibrosis in tumor tissue is considered a histological feature associated with poor prognosis in mammary neoplasms. Although there are other histopathological markers for malignant tumors, it has been suggested that the presence of fibrosis in carcinomas could be used as a prognostic factor in mammary tumors, as it originates from the overreaction of the tumoral stroma, which is comprised mainly of fibroblasts and collagen.

A positive correlation has been reported between fibrosis and mammary tumors in rats when using Picrosirius Red stain, which suggests that neoplasms induced by NMU (N-Methyl-N-Nitrosurea) also have high amount of fibrosis, reinforcing the clinical relevance of fibrosis (McPhail and Robinson, 2010). Furthermore, it has been shown that invasive ductal mammary carcinomas containing fibrosis are more aggressive than those without fibrosis (Hasebe *et al.*, 1996).

In agreement with Soler *et al.* (2016), there were fewer benign tumor samples in the present study than malignant ones, which might explain the lack of significant difference in the parameters between the two types of tumors. Heterogeneity of canine mammary tumors and the limited number of samples make data interpretation challenging; larger samples could result in more significant results. Nevertheless, a positive correlation was observed between histological grade and fibrosis percentage in the tumoral mass, corroborating with the findings by McPhail and Robinson (2010) in rats.

Histologically, fibroblasts and collagen fibers within a fibrotic area are arranged in irregular patterns, associated with increased cellularity and/or fibroblast collagenization. It is still unclear how this dense tissue architecture affects the tumor; however, non-invasive imaging techniques can evaluate collagen *in vivo* and could identify and delineate a fibrotic focus within the mammary tumor (Brown *et al.*, 2003).

Invasive carcinomas with fibrotic foci are more aggressive than those without fibrosis. Fibrosis has been associated with large tumoral size, great histopathological grade, high proliferative activity of tumoral cells, and necrosis. Smaller studies have shown the independent prognostic significance of fibrosis in patients with invasive breast cancer. Furthermore, it has been shown that the relationship between fibrosis and tumor size could be an important prognostic factor (Van den Eyden *et al.*, 2007).

Thus, it has been demonstrated that presence of fibrosis indicates differences in the malignancy of invasive carcinomas and other mammary tumors (Hasebe et al., 2000). Fibrosis is part of the tumor stroma, indicating that the malignant potential of neoplasms depends not only on the biological features of tumoral cells, but also on the tumoral stroma. Fibrosis consists of fibroblasts or collagen fibers mixed with microvessels and the interaction of these components influences the progression of mammary carcinomas with fibrosis (Hasebe et al., 2000). The presence of atypical fibroblasts in the tumoral stroma, especially in the fibrotic foci, is significantly associated with tumor recurrence and death of patients with invasive ductal mammary carcinoma (Ahn et al., 2012).

The influence of an overactive stroma on neoplastic cells, and vice versa, is still unclear. The way in which *in situ* carcinomas remain localized within the basement membrane or become invasive is not fully understood (Hanahan and Weinberg, 2000; Kalliuri, 2003). Furthermore, stromal fibroblasts secrete metalloproteinase (MMPs) and other cytokines that are important factors in tumor progression (Hasebe *et al.*, 2000). Therefore, it is presumed that the biological factors of the usual fibroblasts are markedly different from those in the fibrotic focus. This indicates heterogeneity in fibroblast characteristics associated with tumoral stroma, as well as neoplastic cells, which may affect patients with mammary carcinomas.

A study using 140 cases of invasive carcinomas in women was performed to determine the prognostic significance of foci of fibrosis (Hasebe *et al.*, 2000), which was evaluated by the following histological parameters: age, histopathological grade, nodular state (absent or present), size, tumor necrosis (present or absent), and adjuvant chemotherapy (none or administered). The overall disease-free survival time was significantly lower in cases with fibrosis, confirming that its presence and extent are useful histological and prognostic parameters in patients with invasive lobular carcinoma as well as invasive ductal carcinomas (Hasebe *et al.*, 2000).

Several techniques and different histopathological and molecular markers are currently available to evaluate aggressiveness, biological behavior, and prognosis of mammary neoplasms in bitches. Most markers reflect specific aspects of neoplastic cell biology; however, oncogenesis is determined by complex interactions between epithelial and stromal tumor cells. Markers that reflect the behavior of these cells are scarce in Veterinary Medicine; therefore, in the present study, easy and reproducible evaluation of fibrotic areas was proposed as a practical histopathological marker for tumoral aggressiveness.

This study provides evidence that Masson's trichrome staining could be used for assessing fibrosis in canine mammary tumors. Further *in vivo* and *in vitro* studies are necessary to elucidate the complex pathophysiological development of fibrosis in canine mammary tumors, as well as its impact on the interactions between the immune system and chronic diseases. The number of samples used could have limited the evaluation of the correlation between fibrosis and malignancy; therefore, further studies involving a greater number of samples, different breeds, age, groups, malignant and benign histological types, and different histological staining techniques are needed.

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Conflicts of interest

The authors declare they have no conflicts of interest with regard to the work presented in this report.

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Masculinization of South American catfish (*Rhamdia quelen*) through dietary administration of 17α-methyltestosterone[¤]

Masculinización del pez Liso (Rhamdia quelen) mediante suministro dietario de 17a-metiltestosterona

Masculinização do Jundiá (Rhamdia guelen) através da administração dietética de 17a-metiltestosterona

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Abstract

Background: Production of monosex cultures of females is desirable in commercial aquaculture of certain species due to their higher growth rate. Ideally, females should be free of exogenous hormones. The initial step to produce hormone-free all-female offspring is masculinization of normal females to create sex reversed animals, called neomales, which are then be bred with normal females obtaining all-female offspring. **Objective:** To masculinize *Rhamdia quelen* fry by adding 17α -methyltestosterone (MT) hormone to the feed. **Methods:** Larvae of *R. quelen* were fed diets supplemented with 60, 80, or 100 mg MT/kg feed for 21 days. A control group was fed the same diet free of MT. At 150 days post-hatching, 30 fish of each treatment group were euthanized to evaluate gonadal changes using histological techniques. **Results:** MT significantly affected the differentiation of female gonads in the 60 and 80 mg MT/kg feed groups. Sex reversal was observed in all MT treatment groups, with 50, 40, and 20% neomales produced with 60, 80, and 100 mg MT/kg feed, respectively. Intersex gonads were observed only in the masculinization treatment groups. Inhibitory effects on gonadal development of females and males were observed at the highest MT doses. **Conclusion:** Dietary administration of MT effectively masculinizes *R. quelen* fry; however, the lowest dose of 60 mg/kg feed is recommended, since higher doses have inhibitory effects on gonadal development in both sexes.

Keywords: histology, fish gonads, monosex culture, neomales, sex reversal.

Resumen

Antecedentes: Debido a su mayor tasa de crecimiento, la producción de peces femeninos monosexo es deseable en acuicultura comercial de determinadas especies. Idealmente, las hembras deben estar libres de

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hormonas exógenas. El paso inicial para generar descendencia femenina libre de hormonas es la masculinización de hembras genéticas para producir animales sexualmente revertidos, llamados neomachos, los cuales se aparean luego con hembras genéticas para producir descendencia femenina. **Objetivo:** Masculinizar larvas de *Rhamdia quelen* con 17 α -metiltestosterona (MT) incorporada en el alimento. **Métodos:** Larvas de *R. quelen* fueron alimentadas con dietas suplementadas con 60, 80 y 100 mg de MT/kg de alimento durante 21 días. Un grupo control recibió la misma dieta, sin MT. A los 150 días pos-eclosión, 30 peces de cada tratamiento fueron eutanasiados para evaluación gonadal mediante técnicas histológicas. **Resultados:** La MT afectó significativamente la diferenciación de las gónadas femeninas en las dosis de 60 y 80 mg MT/kg de alimento. El cambio de sexo se observó en los tratamientos con las dosis de 60, 80 y 100 mg MT/kg alimento, con 50, 40 y 20% de neomachos, respectivamente. En los tratamientos de masculinización se observaron gónadas intersexuales. En las dosis más altas de MT se observó inhibición del desarrollo gonadal de hembras y machos. **Conclusiones:** El suministro dietario de MT masculinizó las larvas de *R. quelen*. Se recomienda usar 60 mg/kg, ya que dosis mayores tienen efectos inhibidores en el desarrollo gonadal para ambos sexos.

Palabras clave: cultivo monosexo, gónadas de peces, histología, neomachos, reversión sexual.

Resumo

Antecedentes: O cultivo monossexo feminino é desejável na aquicultura comercial de determinadas espécies devido à maior taxa de crescimento. Idealmente, as fêmeas devem ser livres de hormônios exógenos. O passo inicial para gerar descendências femininas livres destes hormônios é a masculinização de fêmeas normais para produzir animais revertidos sexualmente, os chamados neomachos. Os neomachos podem então ser cruzados com fêmeas normais para a produção de descendências femininas. Objetivo: Masculinizar larvas de R. quelen utilizando o hormônio 17α-metiltestosterona (MT) incorporado no alimento. Métodos: Larvas de R. quelen foram alimentadas com dietas suplementadas com MT nas doses de 60, 80 ou 100 mg/kg de alimento durante 21 dias. Um grupo controle foi alimentado com dieta similar livre de MT. Aos 150 dias pós-eclosão, 30 peixes de cada grupo de tratamento foram eutanasiados para avaliação gonadal através de técnicas histológicas. Resultados: O uso de MT afetou significativamente a diferenciação das gônadas femininas nas doses 60 e 80 mg/kg de alimento. A reversão sexual foi observada em todos os grupos tratados com MT, gerando 50, 40 e 20% de neomachos nas doses 60, 80 e 100 mg MT/kg de alimento, respectivamente. Gônadas intersexuais foram observadas somente nos tratamentos masculinizantes. Nas maiores doses de MT, efeitos inibitórios de desenvolvimento gonadal foram observados em fêmeas e machos. Conclusão: A administração dietética de MT efetivamente masculinizou larvas de R. quelen. No entanto, a dose mais baixa de 60 mg/kg de alimento é recomendada devido aos efeitos inibitórios das doses mais elevadas de MT no desenvolvimento gonadal de ambos os sexos.

Palavras-chave: cultivo monossexo, gônadas de peixe, histologia, neomachos, reversão sexual.

Introduction

Production of monosex fish is highly desirable for many species of commercial interest, such as *Oreochromis* and some salmonids, mainly due to size differences between sexes at maturity or to undesired reproduction during the production cycle (Dunhan, 2004). Monosex fish populations are typically obtained by sex reversal, which can be induced by temperature (Baroiller *et al.*, 1999) or by hormonal administration. Hormones can be applied via the diet, by bath immersion, and injections or implants (Pandian, 2013). Dietary supplementation with hormones is a low-cost method for sex reversal, and is widely used in species with a labile period (period in which sex cells are totipotential) after the first feeding (Pandian, 2013). However, the use of hormones, such as steroids, is controversial for human and environmental health, and many European countries have banned it for farmed fish (Desprez *et al.*, 2003; El-Sayed, 2006). Consequently, monosex populations may also be induced indirectly by the combined use of hormones and mating strategies. This two-stage method of sex reversal has been implemented for the commercial culture of fish, such as salmonids (Hunter *et al.*, 1983; Feist *et al.*, 1995), to avoid direct application of hormones to species raised for human consumption.

The stages to produce hormone-free female offspring involve using hormones to directly masculinize females, followed by performance tests to identify masculinized females, known as neomales.