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Overexpression of 11β-hydroxysteroid dehydrogenase 1 in visceral adipose tissue and underexpression of endothelial nitric oxide synthase in the adrenal cortex of dogs with hyperadrenocorticism

Diego Daniel Miceli^{1,2}, Adriana María Belén Abiuso², Patricia Noemi Vidal¹, María Florencia Gallelli¹, Omar Pedro Pignataro² and Victor Alejandro Castillo^{1,*}

¹Universidad de Buenos Aires, Facultad de Ciencias Veterinarias, Cátedra de Clínica Médica de Pequeños Animales y Hospital Escuela de Medicina Veterinaria, Unidad de Endocrinología, Av. Chorroarín 280, Buenos Aires CP 1427, Argentina

²Laboratorio de Endocrinología Molecular y Transducción de Señales, Instituto de Biología y Medicina Experimental – CONICET, Vuelta de Obligado 2490, Buenos Aires CP 1428, Argentina

Abstract

11 β -Hydroxysteroid dehydrogenase 1 (11 β -HSD1) is an enzyme that activates cortisone into cortisol in tissues. Alterations in this enzyme are related to the development of metabolic syndrome, obesity and hyperadrenocorticism (HAC). Endothelial nitric oxide synthase (eNOS) produces nitric oxide and is related to the regulation of adrenal steroidogenesis. The aim of the study was to evaluate 11 β -HSD1 and eNOS expression in dogs with HAC. Visceral adipose tissue samples were taken to evaluate 11 β -HSD1 expression by immunohistochemistry and western blotting. In parallel, adrenal gland samples were collected to evaluate eNOS expression by immunohistochemistry. 11 β -HSD1 expression was significantly higher in the adipocytes of dogs with HAC than in those of the control dogs. eNOS expression in the adrenal cortex (zona fasciculata) was significantly lower in the dogs with HAC than in the control dogs. 11 β -HSD1 overexpression and eNOS underexpression could play a role in the maintenance of hypercortisolism in dogs with HAC.

Keywords: 11β-HSD1, eNOS, Glucocorticoids, Hyperadrenocorticism.

Introduction

The actions of glucocorticoids in target tissues are determined by the "nuclear" density of their receptors and the intracellular metabolism of the 11βhydroxysteroid dehydrogenase (11B-HSD) isoforms (Chapman et al., 2013). 11β-HSD is an enzyme that interconverts inactive glucocorticoid compounds (cortisone) into active compounds (cortisol) and vice versa (Seckl and Walker, 2001; Sieber-Ruckstuhl et al., 2007). Two isoforms of the enzyme have been described: 11B-HSD1 mainly catalyses the activation of cortisone to cortisol to increase the intracellular cortisol levels and thus amplifies its cellular actions (Morton and Seckl, 2008), whereas 11B-HSD2 catalyses the reverse reaction (Tomlinson et al., 2004). Given the physiological relevance of both 11β-HSD isoforms, they are referred to as "intracellular gate-keepers of tissue glucocorticoid action" (Chapman et al., 2013).

11 β -HSD1 is constitutively expressed in a wide range of tissues, including the liver, adipose tissue, pancreatic islets, bone, gonads, and the nervous system. Additionally, expression of this enzyme can be induced in fibroblasts, smooth and skeletal muscle, and cells of the immune system (Sieber-Ruckstuhl *et al.*, 2007; Ryan *et al.*, 2011; Champan *et al.*, 2013).

Recently, studies on the expression and actions of the 11β-HSD1 enzyme have become relevant because alterations of the enzyme are associated with metabolic syndrome, obesity (Wake and Walker, 2004; Walker and Andrews, 2006), and the metabolic complications of hyperadrenocorticism (HAC) (Morgan et al., 2016). Various studies have reported that 11B-HSD1 expression is increased in the visceral adipose tissues and livers of obese patients (Livingstone et al., 2000; Paulmyer-Lacroix et al., 2002; Candia et al., 2012). This increase in turn increases the local and/or systemic concentrations, thereby favouring its cortisol detrimental effects (Seckl and Walker, 2001; Stulnig and Waldhausl, 2004).

The hypercortisolism of HAC induces a state of insulin resistance that provokes a tendency towards fasting hyperglycemia, hypercholesterolemia, hypertriglyceridemia, abdominal obesity, hepatic steatosis and diabetes mellitus in the most severe cases (Pivonello *et al.*, 2010; Di Dalmazi *et al.*, 2012; Miceli *et al.*, 2014, 2017). There is evidence that local metabolism of glucocorticoids mediated by the 11 β -HSD1 enzyme plays a central role in the development of these complications, which challenges the classical view that active glucocorticoids in general circulation

*Corresponding Author: Victor Alejandro Castillo. Department of Companion Animals, Chief of Endocrinology Unit, Faculty of Veterinary Sciences, University of Buenos Aires, Buenos Aires, Argentina. Email: <u>vcastill@fvet.uba.ar</u>

represent the most important mode of action of glucocorticoids (Morgan *et al.*, 2016).

Nitric oxide (NO) is a gaseous inorganic free radical whose function as an intracellular messenger has been widely established in a variety of cell types (Nathan, 1992). NO is generated by the oxidation of L-arginine in a reaction catalysed by NO synthase (NOS) (Palmer et al., 1988). NO is involved in numerous biological processes, including smooth muscle relaxation, neurotransmission, cytotoxicity, and nociception (Furchgott and Zawadaki, 1980; Palmer et al., 1987; Garthwaite, 1993; Paradise et al. 2010). Three NOS isoforms have been identified: neuronal (nNOS), endothelial (eNOS), and inducible (iNOS) (Sessa, 1994). eNOS is located in many tissues mainly on the cellular plasma membrane but is also found in the cytosol, nucleus, Golgi apparatus, and mitochondria (Oess et al., 2006).

NO is involved in the modulation of various endocrine systems (Van Voorhis *et al.*, 1994; Del Punta *et al.*, 1996; Natarajan *et al.*, 2002). Specifically, in the adrenal glands are known to modulate steroidogenesis in rat adrenal zona fasciculata cells (Cymeryng *et al.*, 1998, 2002), in which an inhibitory effect of NO has been demonstrated. Although adrenal steroidogenesis is regulated primarily by adrenocorticotropic hormone (ACTH), the various cell types that compose the structure of the cortex and adrenal medulla secrete compounds that play roles in the fine-tuning of the regulation of adrenal physiology (Hochol *et al.*, 2004; Conconi *et al.*, 2006; Bornstein *et al.*, 2008; Ducsay and Myers, 2011).

Cortisol decreases the plasma NO levels by inhibiting eNOS (Yang and Zhang, 2004; Liu *et al.*, 2009). In dogs with HAC, numerous systemic and metabolic disturbances are generated when the vasodilatory effect of NO is lost, including hypertension, thromboembolism, and blindness (Loscalzo, 1995; Mitchell and Webb, 2002; Luchi *et al.*, 2003; Cabrera Blatter *et al.*, 2012).

The objective of the present study was to evaluate 11β -HSD1 enzyme expression in visceral adipose tissues and eNOS expression in the adrenal fascicular cortices of dogs with HAC.

Materials and methods

Detection of 11β -HSD1 expression in visceral adipose tissues

Study material

Visceral adipose tissues (n = 16): samples from dogs with pituitary-dependent hyperadrenocorticism (PDH) (n = 8) were obtained by necropsy from patients with PDH who required euthanasia or died for another causes (4 females and 4 males; median age of 10 years, range 9 to 14 years; 3 Crossbreeds, 2 Poodles, 1 Boxer, Labrador Retriever and Dachshund; body condition between 7/9 and 8/9, according to Jeusette *et al.* (2010).

The diagnosis of PDH was based on history, physical examination, routine laboratory findings and an increased urine cortisol:creatinine ratio in two consecutive morning urine samples collected by the owner at home. After collection of the second urine sample, three doses of dexamethasone (0.1 mg/kg) were administered PO at 8-hour intervals, and a third urine sample was collected the following morning, according to Kooistra and Galac (2012). The diagnosis was further supported by visualisation of the bilateral adrenal hyperplasia by ultrasonography and the pituitary mass by nuclear magnetic resonance imaging. Dogs with an infectious disease (systemic or local), pancreatitis. diabetes mellitus, hypothyroidism, tumours (except pituitary adenoma) or any other systemic disorder capable of inducing insulin resistance, or increasing TNF- α , were excluded (Hess, 2010). Also, dog with abnormalities on complete blood count, urinalysis or biochemical profile other than those expected with PDH (stress leukogram, low urine specific gravity, mild hyperglycemia, dyslipidemia, increased alkaline phosphatase or alanine transferase), were excluded. In all cases, the common therapy used in Argentina was administered, consisting of combined treatment with cabergoline (0.07 mg/kg/week, divided into doses administered every 48 hours) and ketoconazole (15-20 mg/kg administered every 24 hours) (Castillo et al., 2008; Lien and Huang, 2008). All the dogs had a good response to the therapy, controlling the clinical signs and regularizing the biochemical alterations in less than 3 months. The dogs with HAC were euthanized (n = 5) when they were seriously compromised with the quality of life of the animal, product of the chronic deterioration of the disease (median survival time: 26 months, range 18 to 35 months).

Samples from control healthy dogs (n = 8) (4 females and 4 males; median age of 7 years, range 5 to 11 years; 4 Crossbreeds, 1 Beagle, Maltese, Yorkshire Terrier; body condition between 5/9 and 6/9) were obtained during minor surgical procedures (e.g., ovariohysterectomy or exploratory laparotomy by foreign body; anesthetic protocol (acepromazinefentanyl as premedication; propofol as induction, insufluorane as maintenance; continuing infusion with remifentanyl). None of these dogs were undergoing any treatment with steroids before sampling.

Immunohistochemical assay

Visceral adipose tissue samples were fixed in 4% buffered formol, embedded in paraffin and processed for histological examination. Slices with a 5- μ m thickness were made from paraffin sections of visceral fat. The sections were deparaffinised by passages in xylol and decreasing alcohol gradients. The sections were washed with PBS (pH 7.6) and permeabilised with 1x Triton X-100 (Sigma-Aldrich, Saint Louis, USA) in

PBS. After washing, peroxidase activity was blocked with 3% hydrogen peroxide (H₂O₂). Further washes were performed, and nonspecific binding sites were blocked with blocking serum (goat serum in PBS) for 10 minutes. Then, the samples were washed with PBS. The sections were incubated for 2 hours at 37 °C with a 1:50 dilution of a rabbit polyclonal antibody against amino acid region 65-164 of the 11B-HSD1 enzyme (Santa Cruz Biotechnology, H-100, sc-20175) in PBS. After two washes, the Immunoperoxidase Detection System (Millipore IHC Select) was used. This system uses the avidin-biotin-peroxidase (ABC) method. Briefly, the sections were incubated for 10 minutes with the biotinvlated anti-rabbit secondary antibody. After another wash, the sections were incubated for 10 minutes with horseradish peroxidase (HRP)-conjugated streptavidin and finally washed with PBS. The reaction was developed with the chromogen 3, 3'diaminobenzidine (DAB). The sections were stained with haematoxylin (as a contrast) for 5 minutes, dehydrated (with increasing alcohol gradients and xylol), and mounted with Canada balsam. In the sections used as negative controls, PBS was included in place of the primary antibody. Image analysis: the DC180 digital camera and Leica trinocular microscope model DMLS were used for image capture. The Qwin Plus software from Leica Inc. Corp. was used. Images of 30 non-consecutive fields were evaluated in each fat slice to determine the intensity of the 11β-HSD1 immunostaining. To this end, two observers (independent from the research project) analysed the images, which were scored according to the intensity of the staining. The following values were assigned (Oberholzer et al., 1996; Ahmad et al., 2011): "0", no immunostaining; "1", weak immunolabelling; "2", moderate immunolabelling; "3". and strong immunostaining.

Western blotting

Visceral adipose tissue samples were frozen immediately at -80°C until processing (10 months). The samples were homogenized in 5 ml of Lysis Buffer (10 mM Tris-HCl [pH 8], 1% Triton X-100, and 0.5 mM acetic acid) with a cocktail of protease inhibitors (25 mM HEPES, 250 mM sucrose, 4 mM EDTA, 1 µM leupeptin, 1 µM aprotinin, and 1 µM pepstatin, pH 7.4). The homogenates were centrifuged at 12,000 x g for 10 minutes at 4 °C. The supernatant was collected, and the protein concentrations were quantified using the Bradford method (Bio-Rad Laboratories) (Sapan et al., 1999). Equal amounts of protein per sample (30-40 g) were boiled for 5 minutes in Lysis Buffer (0.06 M Tris-HCl [pH 6.8], 25% v/v glycerol, 2% w/v SDS, 0.01% bromophenol blue, w/v and 5% v/vβmercaptoethanol). Then, the proteins were subjected to polyacrylamide gel electrophoresis under denaturing conditions (SDS-PAGE) and transferred to a

nitrocellulose membrane (Bio-Rad Laboratories). The membranes were blocked with 5% skim milk in 0.1% PBS-Tween (PBS-T) and incubated overnight (16 hours at 4 °C, constant agitation) with a rabbit polyclonal antibody against the 65-164 amino acid region of the 11β-HSD1 enzyme (Santa Cruz Biotechnology, H-100, sc-20175;1:200 dilution) and the G9545 rabbit anti-GAPDH antibody (1:100,000 dilution). After three washes (10 minutes at room temperature) in PBS-T, the membranes were incubated for 2 hours with the HRP-conjugated anti-rabbit IgG secondary antibody (1:4000 dilution). Finally, the reaction was developed by incubating the membrane for 1 minute in a peroxidase substrate solution for enhanced chemiluminescence (ECL). The bands were digitized using the GBOX and measured with the ImageJ software (NIH, Bethesda, MD, USA). The results are normalized to the expression of a constitutive protein (GAPDH).

Detection of eNOS expression in the adrenal cortex Study material

Adrenal glands (n = 18): samples from dogs with HAC (n = 8) were obtained by necropsy of patients with PDH (the same dogs previously mentioned), whereas control samples (n = 10) (5 females and 5 males; median age of 9 years, range 6 to 14 years; 5 Crossbreeds, 1 Golden Retriever, Beagle, Poodle, Pit Bull Terrier, Jack Russell; body condition between 5/9 and 7/9) were obtained by necropsy of animals euthanized for humanitarian reasons and by veterinary indication (medullary compression and polytrauma). These dogs were euthanized because they arrived at the hospital with a severe trauma, from which there was no possibility of recovery (traffic accident or fall height).

Immunohistochemistry assay

The same immunohistochemistry protocol described above was used. The sections were incubated for 2 hours at 37 °C with a 1:250 dilution of a rabbit polyclonal antibody against eNOS (Thermo Fisher Scientific, PA3-031A) in PBS. *Image analysis:* Images of 50 non-consecutive fields were evaluated in each adrenal cortex to determine the optical density of the cortex (zona fasciculata). The optical intensity analysis was performed after transforming the images to greyscale (8-bit). Calibration for the optical density was performed using a calibrated scale tablet ("step tablet"). The calibration curve ranged from 0 to 2.6 (optical density units) according to Oberholzer *et al.* (1996). Thus, the optical density value increased as the staining strength increased.

Statistical analysis

Prior to analysing the normality of the variables (D'Agostino-Pearson omnibus normality test), the experimental results were analysed using a t test (for unpaired samples, significant at P<0.05) and expressed as the mean \pm SEM. To analyse 11β-HSD1 protein

expression based on the western blotting results, oneway ANOVA and Tukey's test were performed (Graph Pad Prism 6, USA-California).

Ethical approval

The Ethics Committee of the Faculty of Veterinary Science (CICUAL) and the Office of Science at the University of Buenos Aires approved the present study according to the laws on experimentation in animals in Argentina and World Health Organization recommendations. Signed consent was obtained from the dogs' owners for participation in the present project.

Results

Visceral adipose tissue from dogs with HAC clearly express11β-HSD1 enzyme more than normal weighted dog (Fig. 1). Expression average score of the 11β-HSD1 enzyme was significantly higher (P=0.02) in the adipocytes of the visceral adipose tissues from the HAC dogs than in those from the control dogs based on the immunohistochemistry analysis (Control: 1.125 ± 0.227; HAC: 2 ± 0.267). Likewise, 11 β -HSD1 enzyme expression (Fig.2) in the adipocytes was significantly higher in the visceral adipose tissues from the dogs with HAC than in those from the control dogs based on the optical density and western blotting analysis (P=0.041). With respect to the eNOS enzyme (Fig.3), immunostaining of the adrenal cells from fasciculata zone showed more expression (antigen concentration) in control dogs than in HAC dogs. The optical density was significantly higher in the control dogs than in the HAC dogs (P=0.008) (Control: 0.824 ± 0.047; HAC: 0.661 ± 0.039).

Discussion

In the study on visceral adipose tissue, we observed that 11β -HSD1 was expressed at higher levels in the adipocytes of dogs with HAC than in those of the control dogs. These results agree with those reported by Ahmad *et al.* (2011), who showed that the visceral adipose tissues of rats treated with dexamethasone had hypertrophy, hyperplasia and greater 11β -HSD1 expression than control rats. Similarly, Morgan *et al.* (2014) demonstrated that glucocorticoid-treated mice not only had greater 11β -HSD1 expression in their adipose and hepatic tissues than control mice but also exhibited increased enzyme activity.

The role of 11 β -HSD1 in the development of HAC was first described by Tomlinson *et al.* (2002), who reported an extremely rare case of a PDH patient with no Cushing phenotype (no hypertension, no muscle atrophy, and no fat redistribution). These authors demonstrated that the patient had a functional defect in the 11 β -HSD1 enzyme that rendered it inactive, which correspondingly decreased the conversion of cortisone to cortisol at the local level. Later, the same enzymatic defect and the same clinical picture were described in a patient with HAC resulting from an adrenocortical nonhypophyseal tumour (Arai *et al.*, 2008).





Fig. 1. Immunostaining for 11 β -HSD1 (**A**) and Average score of 11 β -HSD1 expression (**B**) in dog visceral adipose tissue. (**A**): Negative control (**a**), healthy normal-weight dogs (**b**), dogs with HAC (**c**). Brown cytoplasm indicates positive labelling, whereas blue cytoplasm and nuclei are negative (haematoxylin). Arrows indicate areas with stronger staining. The scale bar is 100 μ m. (**B**): Average score from HAC dogs are significantly higher (* P=0.02) than in control dogs. Values expressed as the mean ± SEM.



Fig. 2. 11 β -HSD1 expression in visceral adipose tissues determined optical density and western blotting in the control dogs and dogs with HAC. The enzyme are highly expressed in HAC (*P<0.05) according to the optical density analyse (top figure) and western blott (bottom figure). Values are expressed as the mean \pm SEM.



Fig. 3. Immunostaining (A) for eNOS in the adrenal cortex (zona fasciculata) and Optical density (B) of the adrenal cells (zona fasciculata). (A): Negative control (a), healthy normal-weight dogs (b), dogs with HAC (c). Brown cytoplasm and nuclei indicate positive staining, whereas blue cytoplasm and nuclei are negative (haematoxylin). Arrows indicate nuclear staining. The scale bars are 50 (top figures) and 20 μ m (bottom figures). (B): Optical Healthy normal-weight dogs present cells with a higher optical density (greater immunostaining) than dogs with HAC (**P<0.01). Values are expressed as mean ±SEM.

Similar results were found in basic research. 11B-HSD1 knock-out HAC model mice (in all tissues) were protected from the adverse effects of cortisol, with no hypertension, no muscle atrophy, no hepatic steatosis, and no increase in visceral fat (Morgan et al., 2014). Additionally, 11B-HSD1 knock-out mice (adipose) were protected against hepatic steatosis, lipolysis in their adipose tissues, and excess plasma free fatty acids but did not fail to present muscular and dermal atrophy. hypertension, and deposition of visceral fat (Morgan et al., 2014). These results suggest that the local metabolism of glucocorticoids mediated by the 11B-HSD1 enzyme plays a decisive role in the development of the complications of hypercortisolism, thereby challenging the classical view that active glucocorticoids in circulation represent the most important mode of action of glucocorticoids (Morgan et al., 2016). Conversely, the local metabolism of glucocorticoids in visceral adipose tissue would play a central but by no means absolute role in the

development of metabolic disturbances of hypercortisolism, especially hepatic and lipid alterations. In agreement with these results, HAC dogs in general and the dogs evaluated in this study in particular present clinical alterations typical of hypercortisolism.

Interestingly, evaluation of 11β -HSD1 enzyme expression in the dogs by western blotting did not reveal the typical 34 kDa and 68 kDa bands found in humans due to the enzyme's dimeric structure (Mariniello *et al.*, 2006). Instead, we found a single band that was 50 kDa in size. Kannisto *et al.* (2004) also reported the existence of this band in human visceral adipose tissue.

Few studies have evaluated 11 β -HSD1 expression in humans with HAC. Mariniello *et al.* (2006) found no significant differences between 11 β -HSD1 expression in visceral adipose tissue from humans with HAC and healthy normal-weight patients. These authors postulated that the lack of an increase in 11 β -HSD1 expression in patients with HAC could be due to downregulation of the enzyme as a result of chronic overstimulation of plasma glucocorticoids.

Most studies in human medicine on obesity, metabolic syndrome, and 11 β -HSD1 have argued that 11 β -HSD1 is overexpressed in adipose tissues in patients with central obesity compared to control patients (Paulmyer-Lacroix *et al.*, 2002; Mariniello *et al.*, 2006; Espíndola-Antunes and Kater 2007; Candia *et al.*, 2012), with increased local production of glucocorticoids that amplifies the effects. Iwasaki *et al.* (2008) postulated that metabolic syndrome would favour an intracellular Cushing state caused by overexpression of the 11 β -HSD1 enzyme.

For future research, it would be interesting to develop new therapies that act specifically on 11 β -HSD1 and thus be able to control hypercortisolism. Likewise, it would be interesting to evaluate if the different therapies already known for HAC act on 11 β -HSD1, since it has been described that trilostane could have an effect on it (Teshima *et al.*, 2014).

Adrenal steroidogenesis is mainly regulated by ACTH. However, numerous factors not related to ACTH play roles in the regulation of the hypothalamus-adrenal-(Bornstein hypophysis axis et al., 2008). Neuropeptides, growth factors, cytokines, opioids, adipokines, bacterial ligands, and other locally produced molecules synergize or antagonize the action of ACTH (Bornstein et al., 2008; Hochol et al., 2004; Conconi et al., 2006). It is well known that NO regulates steroidogenesis in cells of the adrenal glomeruli zone (Natarajan et al., 2002), besides acting on granulosa-luteal cells of the human ovary (Van Voorhis et al., 1994) and on Leydig cells (Del Punta et al. 1996; Mondillo et al., 2009). Thus, it is postulated that dissociation between ACTH and the plasma glucocorticoid levels would occur under certain circumstances, indicating that the adrenal glands would have some "autonomy" with respect to pituitary regulation (Bornstein *et al.*, 2008). This dissociation would occur under both physiological (Choi *et al.*, 2007) and pathological conditions, including inflammatory processes, critical illnesses, and mental disorders (Vermes and Beishuizen, 2001; Carroll *et al.*, 2007).

In the study on adrenal glands, we observed that the expression of the eNOS enzyme evaluated through immunohistochemistry was lower in the dogs with HAC than in the control dogs. The role of NOS in modulating the synthesis of cortisol in the adrenals has been documented and is known to be involved in the fine regulation (inhibitory) of steroidogenesis (Cymeryng et al., 1998, 2002; Repetto et al., 2010) as an intracellular autoregulatory mechanism. However, no studies have been performed in dogs with HAC for comparison with our observations. Our results support the hypothesis that the lower eNOS expression in the adrenal glands of HAC dogs could reduce the modulatory effect of NO on steroidogenesis, leading to insufficient inhibition of cortisol synthesis. On the other hand, it is well know that cortisol inhibits NOS leading to a decrease in NO circulating (Yang and Zhang, 2004; Liu et al., 2009; Cabrera Blatter et al., 2012). Taking this mechanism into account, hypercortisolism could reduce the expression of adrenal eNOS, and therefore, its inhibitory mechanism of steroidogenesis would be reduced.

NOS is an enzyme that has numerous cellular locations (cytosol, mitochondria, Golgi apparatus, and nucleus) that confer different functions (Oess et al., 2006). In contrast to the observations in the zona fasciculata of the adrenals of the control dogs, where immunostaining indicated strong eNOS expression in the cytoplasm and nucleus, cytoplasmic labelling and, to a lesser extent, nuclear staining was observed in the dogs with HAC. Taken together, these results not only suggest that eNOS is expressed at lower levels in the adrenals of dogs with HAC but also that the expression sites may be reduced. Cymeryng et al. (1998, 2002) showed that eNOS expression was increased in mouse adrenocortical tumours (Y1 line) relative to normal adrenals. Therefore, evaluating eNOS expression in dogs with adrenal tumours producing HAC and the expression of the different isoenzymes in other tissues from dogs with HAC will be interesting.

In conclusion, expression of the 11β -HSD1 enzyme is increased in the visceral adipose tissues of dogs with HAC, which could favour a sustained increase in plasma cortisol in these patients. Similarly, the expression of the eNOS enzyme is decreased in the adrenal glands (zona fasciculata) of dogs with HAC, which could lead to a lack of inhibition of NO-mediated steroidogenesis. Both enzymatic alterations could intervene in the maintenance of hypercortisolism, thereby favouring the establishment and/or maintenance of the comorbidities associated with this disease.

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Conflict of interest

The authors declare that there is no conflict of interests.

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