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ARTERIAL ARRANGEMENT OF THE DORSAL SURFACE OF THE THORACOLUMBAR SPINAL CORD IN RABBITS

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ABSTRACT

The aim of this study was to describe the arterial blood supply of the dorsal surface of the thoracolumbar spinal cord in the rabbit. The study was carried out on 10 adult New Zealand white rabbits. The arterial system of the thoracolumbar spinal cord was injected by using Batson's corrosion casting kit No. 17. On the dorsal surface of the spinal cord, we found two irregular longitudinal arteries receiving dorsal branches of *rami spinales* in 70 % of the cases examined. When present, they were located in the *sulcus lateralis dorsalis* bilaterally. In 20 % of the cases, no longitudinal arteries were found on the dorsal surface. In 10 % of the cases, three irregular longitudinal arteries were present which received dorsal branches of the *rami spinales*. A third of these arteries were located in the *sulcus medianus dorsalis*. Along the entire thoracic and lumbar spinal regions, the dorsal branches were left-sided in 56.5 % of the cases and right-sided in 43.5 %. The rabbit is often used as an experimental model for the study of spinal cord injury. According to the high degree of irregularity or the absence of the arteries in the thoracic section, it can be concluded that there is a high risk of possible ischemic damage in the thoracic part of the rabbit spinal cord.

Key words: dorsal branch; dorsal spinal artery; rabbit; spinal cord

INTRODUCTION

The most frequent adversely affected part of the human spinal cord is the thoracolumbar segment [11]. Injury has been known to occur during various surgical procedures in this particular segment [15]. Although numerous studies dealing with the arterial blood supply of the spinal cord have been published, complications still occur due to spinal cord injury, leading to fatal consequences, such as paraplegia or paraparesis [5]. The complications caused by the spinal cord injury have been observed in several animals [2], [3], [9].

The rabbit is often used as an experimental model for the study of spinal cord injury. We focused on spinal arteries supplying the dorsal part of the spinal cord in the thoracolumbar area.

MATERIALS AND METHODS

The study was carried out on 10 adult rabbits (age 140 days). We used New Zealand white rabbits (breed HY+) of both sexes (female n=5; male n=5) with an average weight of 2.5—3 kg in an accredited experimental laboratory of the University of Veterinary Medicine and Pharmacy in Košice. The animals were kept in cages under standard conditions (temperature 15—20 °C, relative humidity 45 %, 12-hour light period), and fed with a granular feed mixture (O-10 NORM TYP). Drinking water was available to all

animals *ad libitum*. The animals were injected intravenously with heparin (50 000 IU.kg⁻¹) 30 min before they were sacrificed by the intravenous injection of embutramide (T-61, 0.3 ml.kg⁻¹).

Immediately after killing, the vascular network was perfused with a physiological solution. During manual injection through the ascending aorta, the right atrium of the heart was opened in order to lower the pressure in the vessels to ensure an optimal injection distribution. Batson's corrosion casting kit No. 17 using a volume of 50 ml (Dione, České Budějovice, Czech Republic) was used as the casting medium. After polymerisation of the medium, 10% formaldehyde was injected into the vertebral canal between the last lumbar vertebra and the sacrum, and between the last cervical and first thoracic vertebra to fix the spinal cord. After 1-week fixation, the vertebral canal was opened by removing the vertebral arches in the thoracic, lumbar and sacral spinal regions. The prepared spinal cord was fixed in 10% formalin. This study was carried under authority decision No. 2647/07-221/5.

RESULTS

The *rami spinales* of the *arteriae intercostales dorsales* and *arteriae lumbales* enter the vertebral canal at the level of intervertebral foramina in association with the respective spinal nerve roots. The *rami spinales* after entering the vertebral canal send to the spinal cord, ventral and dorsal branches. The ventral branches enter the ventral spinal artery. On the dorsal surface, we found two irregular longitudinal arteries receiving dorsal branches of the *rami spinales* in 70% of the cases (Fig. 1). If they were present, they were located in the *sulcus lateralis dorsalis* bilaterally. In 20% of the cases, no longitudinal arteries were present on the dorsal surface (Fig. 2). In 10% of the cases, three irregular longitudinal arteries were present, receiving dorsal branches of the *rami spinales*. A third of these arteries was located in the *sulcus medianus dorsalis* (Fig. 3). The frequency of the occurrence of individual dorsal branches is shown in Table 1. When the two irregular dorsal spinal arteries are present, they are formed only by fusion of the small cranial and caudal branches arising from the dorsal branches. They formed irregular loops between each other on the same and on the opposite sides. Among the dorsal branches observed in the thoracic region, 60.5% of them were in the left side and 39.5% in the right side. In the lumbar region, the observed dorsal branches were left-sided in 52.5% of cases and right-sided in 47.5% of the cases. Along the entire thoracic and lumbar spinal regions, the branches were left-sided in 56.5% of the cases and right-sided in 43.5% of the cases; this may be related to the left-sided localization of the aorta.

DISCUSSION

Until now, published works dealing with the arterial supply of the spinal cord in the rabbit were concentrated only on the ventral branches entering the *arteria spinalis ventralis* [10, 16]. The dorsal branches of the *rami spinales* enter the *arteriae spinalis dorsales* if they are present in a large num-

Table 1. Frequency of occurrence of the dorsal branches of arterial spinal branches in the thoracolumbar region of the spinal cord

| Level | Frequency of occurrence of arterial spinal branches (%) | |
|-------|---|------|
| | Right | Left |
| Th 1 | 40 | 50 |
| Th 2 | 70 | 40 |
| Th 3 | 50 | 70 |
| Th 4 | 70 | 90 |
| Th 5 | 40 | 100 |
| Th 6 | 30 | 60 |
| Th 7 | 100 | 60 |
| Th 8 | 60 | 100 |
| Th 9 | 0 | 70 |
| Th 10 | 60 | 20 |
| Th 11 | 50 | 80 |
| Th 12 | 0 | 80 |
| Th 13 | 30 | 100 |
| L 1 | 50 | 40 |
| L 2 | 40 | 80 |
| L 3 | 60 | 60 |
| L 4 | 30 | 40 |
| L 5 | 50 | 50 |
| L 6 | 60 | 50 |

L – lumbar segment of the spinal cord;
Th – thoracic segment of the spinal cord

ber. On the left side also, they occurred in a large number. These branches supplied the corresponding sections of the dorsal surface of the rabbit spinal cord. Therefore, the rabbit is an appropriate animal for experimentally induced spinal cord ischemia, and experiments with the rabbit are helpful to prevent the occurrence of spinal cord ischaemia in man. It is concluded that the high irregularity or the absence of the segmental arteries in the thoracic section, causes high risk of ischaemic damage in the thoracic part of the rabbit spinal cord.

In comparison with other laboratory animals, the rabbit spinal cord segment is each supplied with one corresponding radicular artery with minimal or no collateral bloodstream

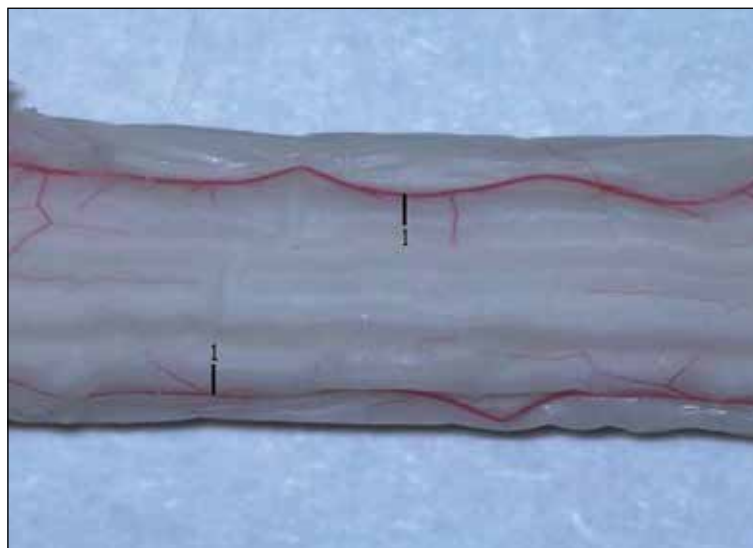


Fig. 1. Presence of two irregular longitudinal dorsal spinal arteries
1 — dorsal spinal artery. Dorsal view. Magn. $\times 12.5$

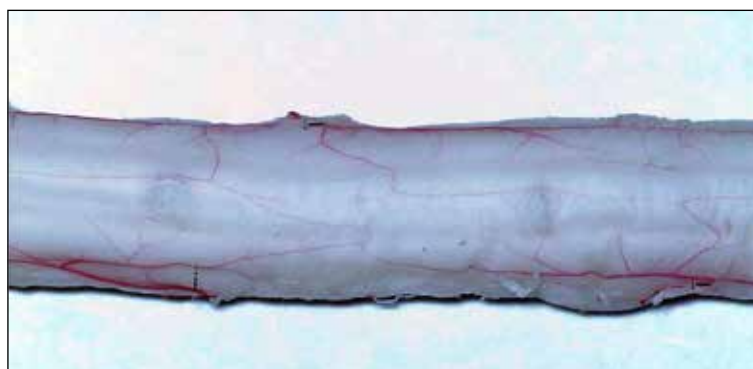


Fig. 2. Absence of longitudinal dorsal spinal arteries
1 — dorsal branch of *ramus spinalis*. Dorsal view. Magn. $\times 12.5$

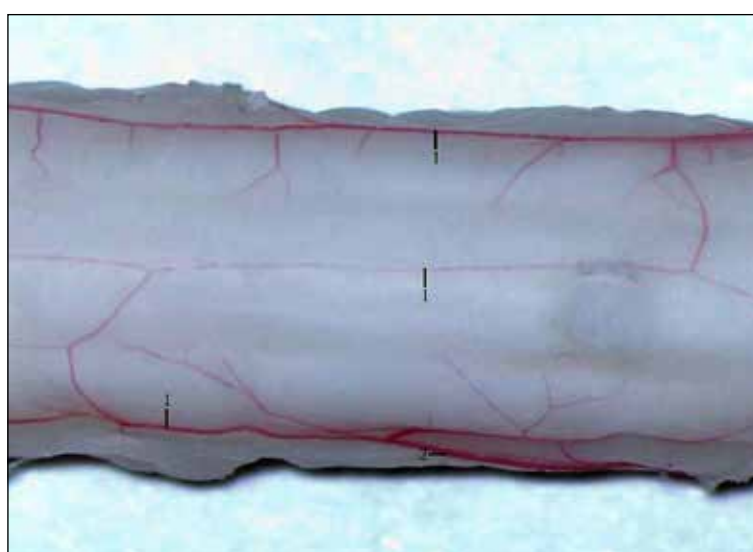


Fig. 3. Dorsal branches of the *rami spinales* forming three irregular longitudinal dorsal spinal arteries
1 — dorsal spinal artery; 2 — dorsal branch of *ramus spinalis*. Dorsal view. Magn. $\times 12.5$

[13]. In the study of the spinal cord ischemic injury, dogs, rats, pigs and mice have been used as experimental models. High variability in the density of the arteries that form the spinal arterial ring and in the *rami spinales* originating from the radicular arteries were found in the dogs [12]. The blood supply of the rat spinal cord has been probably most profusely documented, but the results have been often very different [4], [14], [16], [18]. Two dorsal spinal arteries in the rat were regularly found [8], but in albino Wistar rats they were described as less constant [18]. In pigs, the variations and the presence of extrasegmental arteries of the spinal cord blood supply have been described [17]. Also, in the mouse, spinal cord blood supply was partially described [7]. The disadvantage of experimental work with rats and mice is the small size of the animal and demanding compliance with the standard experimental conditions, such as blood pressure and temperature [13].

Either no or two to three irregular dorsal spinal arteries were located on the dorsal surface (in humans the posterior spinal arteries). The posterior spinal arteries in humans are normally continuous cranial to caudal and supply the posterior third of the spinal cord [6]. In dogs, a pair of dorsal spinal arteries on each side of the dorsal surface of the spinal cord has been described. Each pair was composed of a larger calibre lateral dorsal spinal artery and a thinner medial dorsal spinal artery [12]. In rats, two (much less constant) dorsal spinal arteries that formed irregular loops between each other were described [18]. The dorsal spinal arteries in mice were found in the number of two in some studies [7] or it was described as a single artery [1].

CONCLUSIONS

Until the spinal cord blood supply in the species of experimental animals is not described in detail, it will be very difficult to determine the appropriate species for experiments in this field. Variations in arterial arrangement can produce biased or erroneous results in studies of spinal cord injuries.

ACKNOWLEDGEMENTS

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SEGMENTAL ARTERIES SUPPLYING THE THORACOLUMBAR SPINAL CORD IN THE RABBIT

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ABSTRACT

The aim of this study was to describe the anatomical variations in segmental branches arising from the descending aorta which supply the thoracolumbar spinal cord in rabbits. The study was carried out on twenty adult New Zealand white rabbits. We prepared corrosion casts of the arterial system of the thoracolumbar spinal cord. Spofacryl was used as a casting medium. There were 12 intercostal arteries and 1 costoabdominal artery. In 70 % of the cases 9 pairs were present, in 20 % of the cases 8 pairs, and in 10 % of the cases 10 pairs were found arising from the thoracic aorta. The paired lumbar arteries were present in 6 pairs in 90 % of the cases and in 5 pairs in 10 % of the cases. We found also variations in the level of the origin of the segmental arteries. Based on our results, we concluded that there is a high variability of the blood supply in the thoracolumbar part of the spinal cord in rabbits. Laboratory animals such as the rabbit are frequently used in studies of spinal cord ischemic damage. Variations in arterial arrangements can produce biased or erroneous results in studies of spinal cord injuries.

Key words: anatomical variation; dorsal intercostal artery; lumbar artery; thoracolumbar spinal cord

INTRODUCTION

Rabbits are often used as experimental models for the study of spinal cord injuries and the thoracolumbar area frequently serves as the site for the study of spinal cord ischemia [2], [8]. Until recently, published studies have dealt only partially with the variability of the spinal cord vascular system in the rabbit [5].

The aim of this study is to describe the arterial supply of the spinal cord of rabbits; particularly in the thoracolumbar region where surgical procedures may be associated with a high risk of serious neurological damage.

MATERIALS AND METHODS

The study was carried out on 20 adult rabbits (age 140 days). We used New Zealand white rabbits (breed HY+) of both sexes (female n = 10; male n = 10) with an average weight of 2.5—3 kg in an accredited experimental laboratory of the University of Veterinary Medicine and Pharmacy in Košice. The animals were kept in cages under standard conditions (temperature 15—20 °C, relative humidity 45 %, 12-hour light period), and fed with a granular feed mixture (O-10 NORM TYP). Drinking water was available for all

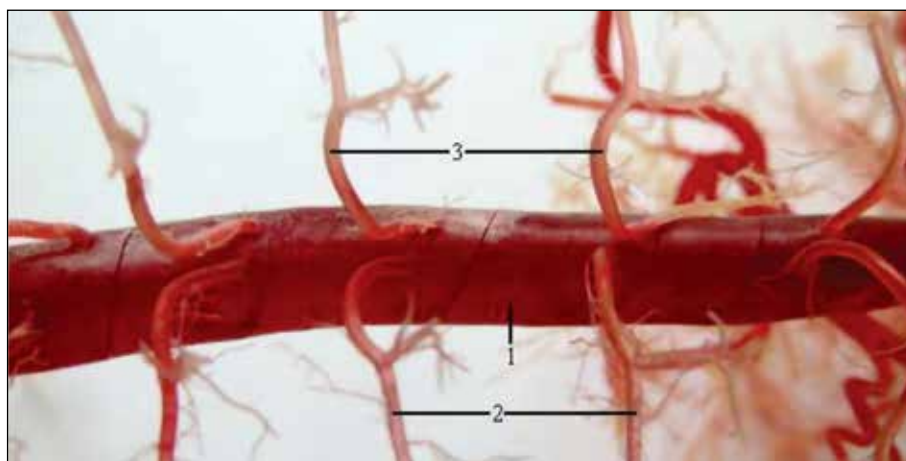


Fig. 1. More cranially located origin of the left sided *arteriae intercostales dorsales* than the right-sided.
(1) *aorta thoracica*; (2) *arteriae intercostales dorsales sinistralae*; (3) *arteriae intercostales dorsales dextrae*.
Macroscopic image, dorsal view



Fig. 2. Origin of *arteriae intercostales dorsales* at the same level as Fig. 1
(1) *aorta thoracica*; (2) *arteriae intercostales dorsales sinistralae*; (3) *arteriae intercostales dorsales dextrae*.
Macroscopic image, dorsal view

animals *ad libitum*. The animals were injected intravenously with heparin (50 000 IU.kg⁻¹) 30 min before they were sacrificed by an intravenous injection of embutramide (T-61, 0.3 ml.kg⁻¹). Immediately after death, the vascular network was perfused with a physiological solution. During the manual injection through the cannula inserted into the ascending aorta *via* the left ventricle, the right atrium of the heart was opened in order to lower the pressure in the vessels to ensure a well perfused injection. Spofacryl (polymethylmethacrylate, SpofaDental, Czech Republic) in a volume of 35 ml was used as the casting medium. The maceration was carried out in 2–4 % KOH solution for a period of 5 days at 60–70 °C. This study was carried out under the authority of decision No. 2647/07-221/5.

RESULTS

The thoracic part of the spinal cord was supplied by *rami spinales*. They were branches from the segmental *arteriae intercostales dorsales*. There were 12 dorsal intercostal arteries and 1 costoabdominal artery (the number of thoracic vertebrae was 13 in all cases). In 70 % of the cases 9 paired dorsal branches arising from the dorsal surface of the *aorta thoracica* were present. In 20 % of the cases 8 pairs, and in 10 % of the cases 10 pairs were present arising from the dorsal surface of the *aorta thoracica*. The remaining first 3–5 pairs branched from the *arteria intercostalis suprema*. In 60 % of the cases, the left-sided arteries originated more cranially

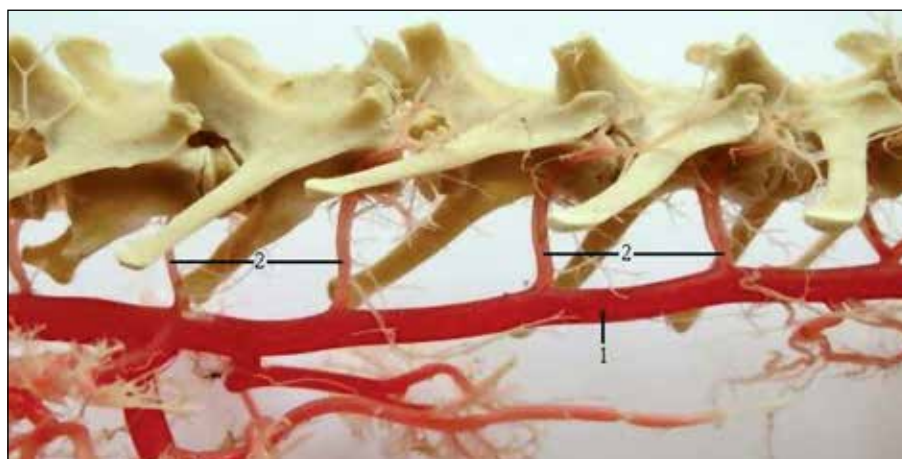


Fig. 3. Origin of *arteriae lumbales* by means of a common trunk
(1) *aorta abdominalis*; (2) *arteriae lumbales*.
Macroscopic image, lateral view



Fig. 4. Origin of *arteriae lumbales* by means of a common trunk. The first two pairs originated as independent branches
(1) *aorta abdominalis*; (2) *arteriae lumbales* with independent origin; (3) *arteriae lumbales* originating by means of a common trunk.
Macroscopic image, lateral view

than the right-sided (Fig. 1). In 20 % of the cases, the right and left-sided arteries originated at the same level (Fig. 2). In 10 % of the cases, the first 9 pairs originated at the same level and in the remaining pairs, the left-sided arteries originated more cranially than the right-sided. In 10 % of the cases, in the first 8 pairs, the right-sided arteries originated more cranially than the left-sided and in the remaining pairs, both the right- and left-sided arteries originated at the same level.

The paired *arteriae lumbales* originated from the dorsal surface of the *aorta abdominalis*. Their *rami spinales* supplied the lumbar spinal cord. They were present in 6 pairs in 90 % of the cases and in 5 pairs in 10 % of the cases (the number of lumbar vertebrae was 6 in all cases). The remain-

ing last pair branched from the *arteria sacralis mediana*. The right and left-sided *arteriae lumbales* originated by means of a common trunk in 60 % of the cases (Fig. 3). In 30 % of the cases, the first 2 pairs were found as independent branches with their origin at the same level (Fig. 4) and the remaining 4 pairs originated by means of a common trunk. In 10 % of the cases, the left-sided arteries of the first 2 pairs originated more cranially than the right-sided and the last 4 pairs originated from the *aorta abdominalis* by means of a common trunk.

DISCUSSION

Similar to other species of farm animals, the blood supply to the thoracic section of rabbit spinal cord occurs *via arteriae intercostales dorsales*. The origin of these arteries branching from the thoracic aorta in laboratory animals varies depending upon the species. The first 3—5 intercostal spaces in rabbits are supplied with blood by *arteriae intercostales dorsales*, which branch off of the *arteria intercostalis suprema*; first 5 in guinea pigs, first 2 in rabbits and first 3 in mice and hamsters [6]. *Rami spinales*, which supply the spinal cord, originate from these arteries. These arteries pass through *foramina intervertebralia* [7]. The remaining portion of the thoracic spinal cord is supplied by the *arteriae intercostales dorsales*, the branches of the *aorta thoracica*. Mice and hamsters have 9 *arteriae intercostales dorsales*, rabbits and hamsters 7 [7] and rats 8 [4]. According to our observations, *arteriae intercostales dorsales* originated irregularly from *aorta thoracica*. Only in 20 % of the cases, do they originate from *aorta thoracica* at the same level. In the majority of corrosive casts, the left *arteriae intercostales dorsales* originated more cranially than the right ones.

The lumbar part of the spinal cord is supplied by means of the *rami spinales* arising from the *arteriae lumbales*. These arteries originate from the *aorta abdominalis*. They are present in 6 pairs in rabbits and in 5 pairs in the rat, guinea pig, mouse and hamster [6]. In these species, the *arteriae lumbales* are present as independent branches arising from the *aorta abdominalis*, but in the rabbit we found the origin of the segmental *arteriae lumbales* at the same level by means of a common trunk.

CONCLUSIONS

The organization with regard to the origin of segmental *arteriae intercostales dorsales* and *arteriae lumbales* has a very important role during operations of thoracoabdominal aneurysms [1]. Correctly performed re-implantation of seg-

mental arteries decreases the risk of spinal cord ischemia, which can also lead to the paraplegia [3]. Until now, only the origin of these segmental arteries as paired branches arising from the dorsal surface of the descending aorta have been described [6], [7].

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LABORATORY DIAGNOSTIC METHODS FOR THE IDENTIFICATION OF *MALASSEZIA* SPECIES A REVIEW

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ABSTRACT

The taxonomy and nomenclature of the genus *Malassezia* was controversial for decades until 1990. Only three species of this genus were known: *M. pachydermatis*, *M. furfur* and *M. sympodialis*. Since the advent of newer molecular methods, new species of *Malassezia* have been discovered. The diagnostics based on phenotypic and genotypic identification of *Malassezia* is the main focus of this article. The phenotypic identification includes: a microscopic survey; cultivation on selective cultivating media; and biochemical examinations. Nowadays, PCR methods, fingerprinting methods, DNA sequence analysis and restriction analysis of PCR amplicons are used for the genotypic identification.

Key words: cultivation; identification; incidence; *Malassezia* spp.; molecular methods; PCR

INTRODUCTION

Yeasts belonging to the genus *Malassezia* are usually members of the normal microflora of the human skin and inhabit the skin of a variety of animal species. All of them are lipophilic yeasts. However, these yeasts are associated with many dermatological disorders of the human skin, such as; atopic dermatitis, dandruff, folliculitis, pityriasis versicolor (PV) or seborrheic dermatitis [31], and intravascular catheter-acquired infections [49]. They have par-

ticipated in different skin disorders in animals, especially otitis externa and dermatitis [23].

Currently, the genus *Malassezia* includes 14 species; *M. globosa*, *M. restricta*, *M. slooffiae*, *M. obtusa*, *M. furfur*, *M. sympodialis*, *M. japonica*, *M. yamatoensis*, *M. dermatis*, *M. pachydermatis*, *M. caprae*, *M. equina*, *M. nana* and *M. cuniculi*, of which, the last four have only been isolated from domestic animals [14]. *M. pachydermatis* is commonly present on the skin and mucosa in dogs and cats but it is not a member of the normal human microbiota. *M. pachydermatis* has also been isolated from ears and from healthy or lesional skin of canines and numerous other animals [42]. *M. pachydermatis* and *M. furfur* can cause life-threatening fungaemia and deep mycoses in immunocompromised patients and preterm neonates [42], [43], [52]. *M. furfur* has been isolated from the scalp, face, dandruff, arms, legs, urine, blood, hair, nails, eyes and the nasal cavity in humans and occasionally from cats, dogs, horses, cows (healthy or with otitis) and bats [7].

M. dermatis has been detected in patients with atopic dermatitis as an associated pathogen of skin lesions [58]. *M. japonica* has been observed from healthy human skin and from the skin of atopic dermatitis patients [59]. *M. obtusa* is a rare species, which is mainly isolated from healthy human skin. The species has occasionally been detected in animals such as dogs with otitis [22] or healthy horses and goats [24]. *M. globosa* has been found in healthy and diseased human skin (PV, seborrheic dermatitis and atopic dermatitis) [21]. This species has occasionally been isolated from animal skin such as cats [9], horses and domestic ruminants [24].

M. restricta occurs mainly on the head, including scalp, neck,

face and ears [3]. *M. yamatoensis* is an infrequent species, which has been reported in patients with atopic or seborrheic dermatitis but seldom from healthy individuals [57]. *M. sympodialis* is an inhabitant of healthy human skin. It occurs especially on the back and chest, but also at other body areas, e.g. the ear canal [54]. *M. sympodialis* has also been isolated from human PV [32] and occasionally from healthy feline skin [20], goats [24] and cows [25]. *M. slooffiae* is found in low frequency on healthy or lesioned human skin, usually in association with *M. sympodialis*, *M. furfur*, *M. globosa* or *M. restricta* [32]. *M. slooffiae* is also commonly isolated from animal skin, from pigs [35], cats [11] cows [25] and goats [61].

M. nana has been found in healthy cats [11], cats with otitis externa, healthy cows or those with otitis [38]. *M. equina* was isolated from healthy anal skin of horses and from the skin of cows in Spain [13]. *M. caprae* so far, has been found on the healthy skin of goats and horses [13]. *M. cuniculi* has been found on healthy skin of the external ear canal of a rabbit in Spain [14].

The main focus of this article is the description of the laboratory diagnostics of the members of the genus *Malassezia*, based on phenotypic and genotypic identification.

IDENTIFICATION OF MALASSEZIA SPP.

The identification of *Malassezia* spp. is complicated by the difficult isolation and cultivation which is needed. Cultivation requirements differ by the type of species [37].

Several approaches have been used to routinely identify *Malassezia* species. These include phenotypic and genotypic identifica-

tion. Phenotypic identification (biochemical and morphological characteristics) (Table 1) include: assessing of the growth on cultivating media; growth with different *Tween* nonionic detergents as the sole lipid supplement; the presence of catalase; temperature requirements [34]; the presence of β -glucosidase revealed by the splitting of esculin; selective growth with cremophor EL; and cell morphology [50]. *Malassezia* cells are unipolar, with size varying from 1.5 to 10 μ m. They are usually round or ovoid in shape and bud on a broad or narrow base [58]. In contrast to other *Malassezia* spp., *M. globosa* cells are typically spherical, with buds growing from a narrow base [3]. Morphologically these species cannot be distinguished, therefore the phenotypic identification is necessary. *M. pachydermatis* is a lipophilic, but not a lipid-dependent specie. It grows on Sabouraud's dextrose agar [10]. The other 13 *Malassezia* species have lipophilic character and therefore many culture media are supplemented with various lipid sources [2]. The following media are the most used; Dixon's agar medium [62], Modified Dixon's agar medium [32], Leeming and Notman agar medium [47], Modified CHROM agar Candida [44], and Modified CHROMO agar [45].

The main physiological tests applied for identification were based on the catalase reaction and the ability of species to utilize different Tweens [20], [40], [60], [80], [34].

Since the recognition of other lipid dependent species, molecular techniques (Table 2) are needed for better diagnostics. For instance, 13.8% of isolates identified by phenotypic means were found to be misidentified after molecular reidentification using sequence analysis of the D1/D2 domains of the large subunit ribosomal rDNA (LSU rDNA) and the ITS1 and ITS2 regions [36].

Table 1. Differential phenotypic diagnostics of various *Malassezia* species, modified according to Cabañes [14]

| Species/ phenotype characteristics | <i>M. furfur</i> | <i>M. obtusa</i> | <i>M. pachydermatis</i> | <i>M. japonica</i> | <i>M. slooffiae</i> | <i>M. globosa</i> | <i>M. yamatoensis</i> | <i>M. restricta</i> | <i>M. nana</i> | <i>M. sympodialis</i> | <i>M. equina</i> | <i>M. caprae</i> | <i>M. dermatitis</i> | <i>M. cuniculi</i> |
|--|------------------|------------------|-------------------------|--------------------|---------------------|-------------------|-----------------------|---------------------|----------------|-----------------------|------------------|------------------|----------------------|--------------------|
| Growth at 37°C | + | -/w | + | + | + | -/w | + | v | + | + | w | -/w | + | + |
| Growth at 40°C | + | - | + | - | + | - | - | - | v | + | - | - | + | + |
| Utilization of Tweens | 20 | + | - | - | +w | - | + | - | +/- | - | - | - | + | - |
| | 40 | + | - | + | w | + | + | - | + | + | w | w/- | + | - |
| | 60 | + | - | + | + | + | + | - | + | + | w | w | + | - |
| | 80 | + | - | + | - | - | + | - | w | + | w | w | + | - |
| Cremophor EL | +/- | - | + | ? | - | - | ? | - | - | -/w | - | - | w/+ | - |
| Catalase reaction | +/- | + | +w | + | + | + | + | - | + | + | + | + | + | + |
| β -GT | -/w | + | +/- | ? | - | - | ? | - | - | + | - | +/- | ? | + |

w — weak; v — variable; ? — not specified; β -GT — β -glucosidase test

Table 2. Molecular methods that have been applied to detect and identify *Malassezia* species [12]

| |
|--|
| Fingerprinting methods |
| PFGE-Pulsed field gel electrophoresis |
| RAPD-Random amplified polymorphic DNA |
| AFLP-Amplified fragment length polymorphism |
| DGGE-Denaturing gradient gel electrophoresis |
| DNA sequence analysis |
| Ribosomal DNA analysis (D1/D2 region LSU-Large subunit — rDNA) |
| Analysis ITS (Internal transcribed spacer regions) rDNA |
| Analysis IGS (Intergenic spacer regions) rDNA |
| Restriction analysis of PCR amplicons |
| RFLP-Restriction fragment length polymorphism |
| tFLP-Terminal fragment length polymorphism |
| Chitin synthase gene sequence analysis |

DESCRIPTION OF THE MOLECULAR METHODS FOR DETECTION OF MALASSEZIA SPECIES

FINGERPRINTING METHODS

Pulsed field gel electrophoresis (PFGE)

Electrophoretic karyotyping using PFGE demonstrates heterogeneity in chromosomal patterns among *Malassezia* yeasts [41]. Analysis of *M. pachydermatis* isolates demonstrated that the karyotypes of *M. pachydermatis* were similar to each other and contained five chromosomes (number of chromosomes: *M. globosa*-8, *M. obtusa*-6, *M. restricta*-9, *M. slooffiae*-7, *M. sympodialis*-7, *M. furfur* type I-7/8, *M. furfur* type II-10/11) [5]. No variation in karyotype was detected within individual species, with the exception of two distinct types for *M. furfur*, rendering the approach useful for identification [4]. Karyotypes of; *M. caprae*, *M. dermatitis*, *M. equina*, *M. japonica*, *M. nana*, *M. cuniculi* and *M. yamatoensis*, have not been analyzed. Another remarkable feature of the *Malassezia* genomes is their small size, ranging from 6.4 to 14 Mb [64]. Although useful, PFGE is time consuming to perform, requires large amounts of high molecular weight genomic DNA for analysis, and is thus less suited as a routine diagnostic tool [16].

Denaturing gradient gel electrophoresis (DGGE)

Using this method, it is possible to perform the separation of the PCR fragments of the same size, but belonging to different species and differing in their sequence. It is based on the differences in mobility of partially denatured double-stranded DNA in a denaturing acrylamide gel [53]. By the DGGE (SSU region), seven *Malassezia* species (*M. furfur*, *M. globosa*, *M. restricta*, *M. sympodialis*, *M. pachydermatis*, *M. slooffiae* and *M. obtusa*) could be distinguished. In addition, no intraspecific variation was observed. This method is useful for the identification of *Malassezia* isolates, but the clinical use may be hindered by technical demands [60]. DGGE is suitable especially for the analysis of the clinical samples that may include more different species [6].

Random amplified polymorphic DNA analysis (RAPD)

RAPD typing of *Malassezia* isolates has been used in various studies. The use of RAPD typing for taxonomic and henceforth

epidemiological purposes has also been investigated [17]. Representatives of all known species at that time were investigated by 20 decamer primers. All species can be discriminated, but a simple analysis was complicated by the presence of significant intraspecific variation [4]. Similar results were obtained in epidemiological studies of isolates from the skin, where *M. sympodialis* showed the greatest homogeneity [19].

In 55 isolates of *M. Pachydermatis*, which were obtained from various domestic animals and body parts, RAPD discriminated four genotypes, one of them occurred in various animals (cat, horse, goat, pig) and another three were observed only in the external ear canal of dogs. Different genotypes could occur on a single animal or even at a single body site [18]. Other studies on *M. pachydermatis* also demonstrated genetic heterogeneity [40]. Duarte and Hamdan [26] showed genetic heterogeneity between RAPD profiles of *M. furfur* and *M. slooffiae* isolates from human and cattle and between *M. pachydermatis* isolates from dogs and cattle, suggesting that molecular differences of *Malassezia* isolates could be caused by new host adaptations. The authors suggested that RAPD is a useful epidemiological tool.

The most distinguished benefits of the RAPD are; flexibility, facility of technical realization, and high separatory capability. This method does not require the figure of nucleotide sequence of DNA target. RAPD-PCR is an efficient, sensitive and suitable mean of distinguishing closely related strains because of low levels of misclassification and high levels of specificity. The importance of the RAPD typing method have been demonstrated in several studies of *Malassezia* genera and other fungi that are poorly characterized allowing a rapid evaluation of genetic diversity of these species [29].

Amplified fragment length polymorphism analysis (AFLP)

AFLP is a widely used multilocus DNA typing method. It is based on the digestion of genomic DNA with two restriction enzymes, ligation of adapters to the ends of the restriction fragments and PCR amplification using selective primers that are complementary to the restriction half sites and are elongated with one to three selective nucleotides. AFLP has been successfully applied to understand the genetic relationships among isolates of *Malassezia* [13]. Using AFLP all species recognized could be separated [8] and in some species distinct genotypes could be differentiated [36]. AFLP genotyping and RAPD have proved to be useful methods for subtyping and discriminating between *Malassezia* strains [60]. This method is not sufficiently reliable for the identification and discrimination of clinically isolated yeasts [51].

DNA SEQUENCE ANALYSIS

Ribosomal DNA sequences

For the identification of fungal microorganisms including clinically relevant yeasts, the ribosomal RNA (rRNA) genes (also referred to as rDNA, but this also includes the IGS regions) have been often used. Fungal rRNA genes occur in tandem repeats, with each repeat encoding 18S (SSU-small subunit), 5.8S, 26(28)S (LSU-large subunit) and 5S rRNA genes (Fig. 1). Two spacer regions exist in each repeat, namely the Internal Transcribed Spacer (ITS) regions located between SSU and LSU and the InterGenic Spacer (IGS) region located between LSU and SSU. The four subunits have

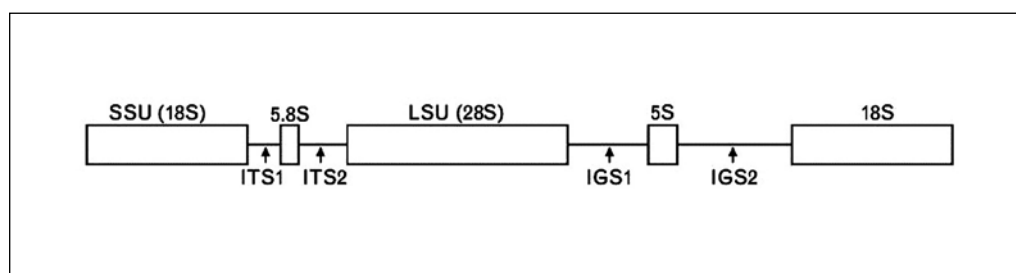


Fig. 1. rDNA gene complex [63]

The rRNA gene complex consists of multiple copies of the small subunit or 18S, the 5.8S region flanked by the ITS regions 1 and 2, the large subunit or 28S and the ITS 1 and 2 that flank the 5S rDNA

more or less constant lengths regardless of the species concerned as follows: SSU — 1800bp, 5.8S — 160bp, LSU — 3500bp and 5S — 120bp. The rRNA genes can be compared among phylogenetically distant species because the sequences are highly conserved, but phylogenetically closely related species or isolates can be compared as well, because of the presence of more variable stretches of DNA, such as the ITS and IGS spacers. At present, sequences of the rRNA genes and ITS regions of almost all human and animal pathogenic fungi have been deposited in the GenBank [7].

DNA sequencing of ITS1 of ribosomal DNA in *Malassezia* spp. was performed to type the species and the strains, because the sequencing is comparatively quick and specific. ITS1 is located between 18S and 5.8S rDNA and is 162–266bp long. Eight species were identified; *M. furfur*, *M. globosa*, *M. restricta*, *M. sympodialis*, *M. pachydermatis*, *M. slooffiae*, *M. obtusa* [48] and *M. nana* [39]. Intraspecific variants have also been discovered within some species (*M. furfur*, *M. sympodialis*, *M. slooffiae* and *M. pachydermatis*) based on sequence variations ranging from 1.7 to 2.8% [48].

Using direct DNA sequencing of LSU of rDNA, the same eight *Malassezia* species were identified [33].

Sequence analysis of IGS1 has not been employed widely for species identification, although sequence variation in this spacer, which is located between LSU and 5S of the rRNA gene, has been employed to classify *M. globosa* and *M. restricta* variants according to their occurrence on seborrhoeic, atopic or healthy skin [55, 56].

RESTRICTION ANALYSIS OF PCR AMPLICONS

Restriction fragment length polymorphism (RFLP)

Enzyme digestion of PCR amplicons has shown to be useful for the differentiation of *Malassezia* species. Primers ITS1, ITS3, ITS4 were used to amplify the ITS region and primers Malup and Maldown were used to amplify the LSU rRNA. PCR amplification of the ITS region with primers ITS1 and ITS4 readily distinguished *M. sympodialis* from other *Malassezia* species by its smaller amplified fragment (700 bp product). The other seven species produced with primers ITS1 and ITS4, the 800 bp product. Five of the seven *Malassezia* species (*M. globosa*, *M. restricta*, *M. slooffiae*, *M. obtusa*, *M. furfur*, *M. sympodialis* and *M. pachydermatis*) could thus be distinguished using the combination of the LSU and ITS regions and restriction endonucleases (LSU with *AvaI* and ITS with *EcoRI* and *NcoI*) [38].

A PCR-RFLP method targeted toward 26S rDNA and with two restriction enzymes, *CfoI* and *BstF51* was developed to identify eleven *Malassezia* species: *M. globosa*, *M. restricta*, *M. slooffiae*, *M. obtusa*, *M. furfur*, *M. sympodialis*, *M. japonica*, *M. yamatoensis*, *M. dermatis*, *M. pachydermatis* and *M. nana* [51].

Distinction of seven *Malassezia* species (*M. furfur*, *M. globosa*, *M. restricta*, *M. sympodialis*, *M. pachydermatis*, *M. obtusa* and *M. slooffiae*) was achievable upon identifying polymorphisms following the digestion of the ITS 3/4 amplicon by *AluI* and *HinfI* [28].

The restriction endonucleases *AluI*, *BanI* and *MspAII* were selected for producing distinct RFLP patterns and were used for the digestion of the PCR products of eleven *Malassezia* species (*M. globosa*, *M. restricta*, *M. slooffiae*, *M. obtusa*, *M. furfur*, *M. sympodialis*, *M. japonica*, *M. yamatoensis*, *M. dermatis*, *M. pachydermatis* and *M. nana*) [27].

Terminal fragment length polymorphism analysis (tFLP)

The tFLP method uses only three different primer sets, minimizing the potential bias related to amplification efficiency. The first set of PCR primers were selectively designed to span the 18S gene through the 28S gene (including ITS1, ITS2 regions and the 5.8S gene) of the fungal rRNA gene. The second two sets of PCR primers were designed to amplify either the ITS1 region or the ITS2 region, in which differences in lengths among *Malassezia* species have been observed. This second step provides additional amplification to increase sensitivity and produces two fragments associated with each *Malassezia* species. The results from this study also show that tFLP analysis is capable of reproducibly assessing the *Malassezia* species present in complex mixtures and human scalp samples. Importantly, it is specific for fungi and is sufficiently sensitive to allow direct assessments of human scalp swab specimens without the need for prior cultivation. Because clinical assessments can be made without prior cultivation, the results are free from culture bias and the turn-around time for analysis is significantly reduced [30].

Chitin synthase gene sequence analysis

Sequencing of the *chs-2* gene has also allowed the differentiation of seven *Malassezia* spp., in spite of the high similarity (95%) in sequence among them [46]. In addition, sequence analysis of *chs-2* indicated that the clinical isolates of *M. pachydermatis* from cats and dogs constituted four distinct genetic types (A, B, C and D), which were linked to skin lesions or otitis [1]. Recently, a mul-

tilocus approach, employing the sequencing of the *chs-2* gene, ITS1 and LSU, has been applied to *M. pachydermatis*. Using this approach, three major *M. pachydermatis* genotypes (A, B and C) were defined [15]. Interestingly, two of these genotypes were associated with skin lesions and high phospholipase activity, whereas the third was isolated from healthy skin and had a low phospholipase activity [17]. Although the multilocus sequencing applied to *M. pachydermatis* provides exciting prospects for population genetic and epidemiological investigations, this approach has not yet been employed for studying *Malassezia* from humans or animals other than dogs [16].

CONCLUSION

At present we know of up to 14 species of *Malassezia* and a large number of strains. These can be divided into three groups according to the host organism: *Malassezia* occurring only in animals (*M. caprae*, *M. equina*, *M. cuniculi* and *M. nana*); *Malassezia* occurring primarily in humans (*M. dermatis*, *M. japonica*, *M. obtusa*, *M. restricta*, *M. yamatoensis*); and *Malassezia* occurring in animals and humans (*M. furfur*, *M. globosa*, *M. slooffiae*, *M. sympodialis* and *M. pachydermatis*). The diagnosis of *Malassezia* comprises phenotypic and genotypic identification. The phenotypic examination is not sufficiently sensitive and therefore the genotypic diagnosis is necessary for exact identification. In the selection of a method for genotypic identification: the technical complexity of a method; diagnostic options without prior cultivation (directly from the samples); time-consuming economic demands; and ultimately the ease of interpretation of the results, play important roles.

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SKIN REPAIR: FROM BIOMECHANICAL, HISTOLOGICAL, AND SPECTROFLUORIMETRIC EVALUATION TO LOW-LEVEL LASER THERAPY OF INCISIONAL AND EXCISIONAL WOUNDS

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ABSTRACT

The use of a simple and reproducible model represents a basic step in an attempt to objectively evaluate the effects of external factors on the wound healing process. In general, there exist two basic models of wound healing: incisional (sutured) and excisional (open). Presented here are our previously published critical review papers of both models of healing which are compared to other studies. Our detailed biomechanical investigations of incisional wound healing, have shown that on the skin regions with minimal strength, there may not be any difference in removing the skin sutures two, three, four, and/or five days following skin closure. Furthermore, an intradermal running suture is more appropriate for experimental use in rats when compared to simple interrupted cutaneous sutures. The spectrofluorimetric studies have shown that fluorescence spectroscopy might be found useful in non-invasive monitoring of selected redox parameters during the early stages of healing and it also allows for the visualization of ischemic areas of the skin. By evaluating the effects of low-level laser therapy (LLLT) on skin wound healing, it was found that lasers modulate open and sutured wound healing differently. The main difference between primary and open/secondary wound healing is in the amount of granulation tissue (GT) formation. Whereas, a primary wound heals without GT formation, a secondary wound needs new tissue development which includes extensive GT formation. Accordingly, we concluded that LLLT should be wound-type (incision vs. excision) specific.

Key words: light therapy; optical diagnostics; tensile strength; wound healing

INTRODUCTION

Due to the expected increase in the occurrence of metabolic, cardiovascular, and oncologic diseases, it may be estimated that the mortality associated with impaired wound healing will increase. In 2003, the health services in the USA spent over 9 billion USD for the treatment of complications related to impaired wound healing [3]. Many factors, such as, diabetes mellitus, AIDS, bad nutrition, as well as chronic application of corticoids, and/or infections, may negatively affect the process of wound repair. In addition, the repair process may differ due to individual biological variations, leading to different outcomes. Stagnation of granulation tissue formation, delayed epithelization, wound dehiscence, creation of hypertrophic and keloid scars, etc., are the most serious complications. Hence, it is necessary to search for new alternatives which either promote wound healing or improve the early diagnostics in post-surgical periods and after traumas. Therefore, a number of experimental studies have dealt with the development of new therapeutic approaches that have the potential to improve skin wound healing, in both healthy and unhealthy individuals [3], [12], [13], [34], [40], [47].

One of many possibilities of how to improve wound healing, is the use of low-level laser therapy (LLLT). LLLT belongs to a group of photochemical interactions of low-energy laser radiation with tissues and cells of living organisms. It has already been shown that

laser radiation affects cell proliferation, differentiation, and the production of the extracellular matrix (ECM). At the present time, however, there is no general agreement about the exact molecular mechanisms of how LLLT influences these biological processes. In numerous clinical and experimental studies it has been proven that low-energy laser light reduces pain, accelerates wound healing and positively influences inflammatory, proliferative, and maturation phases [22], [26], [35], [43], [48]. On the other hand, it has also been shown that the healing after LLLT was not accelerated [8], [28], [29], [41], which contributed to a general skepticism and prevents its common use in clinical practice.

Basic models of wound healing

The use of a simple and reproducible model is a basic requirement for an objective statement of the effects of different external factors on wound repair [18]. In general, there exist two basic models (Fig. 1) of skin wound healing, i.e. incisional and excisional [10]. Whereas the incisional (sutured) skin healing model is preferred for wound tensile strength measurement [9], the excisional (open) model is more appropriate for histological, biomechanical, and molecular evaluation due to a significantly higher amount of new tissue that is formed during the healing process.

On the other hand, in the case of skin incisions, the selection of the appropriate suturing technique represents the principal question. Traditional skin suture techniques are the most commonly used methods of wound closure worldwide. Techniques which appose wound edges evenly, result in a more rapid repair process and better cosmetic outcome. These methods have been proven to be safe and effective. In general, simple interrupted percutaneous suture knotting is the most frequently used method of wound closure [1], [38], [47]. However, continuous intradermal running suture has many advantages in experimental, pediatric, plastic, and veterinary surgery. Intradermal suture reduces the risk of self-induced trauma, since there is no protruding material for the patient to lick or scratch and may be found more appropriate for the experimental use in rats [15]. Of note, the numbers of stitches, their placement and tying represent the main limitation of this wound closure technique.

Rat skin as A model to study wound REPAIR

The rat skin represents one of the most common models used in experimental studies concerning wound healing. Since the rat skin contains three layers such as epidermis, dermis, and striated muscle (*panniculus carnosus*) (Fig. 2), it is a useful model to study the healing of three different tissue types. Nevertheless, most published papers evaluate the effect of treatment only on the epidermis and dermis. The omission of the muscle layer is; however, understandable due to its absence in humans and other frequently used animal models, such as porcine skin.

Only the epidermis is able to completely regenerate. A wound healing of the injured dermis runs in three phases, such as, inflammation, proliferation, and maturation [4]. These phases are not strictly separated from each other, their processes freely blend together. The inflammatory phase is characteristic by a demarcation line formation, consisting mainly of neutrophils, that separates necrotic from vital tissue. During the proliferation phase, fibroblasts and endothelial cells form granulation tissue. Finally, the newly formed tissue loses cells and the remaining collagen becomes organized and forms a scar.

In the healing process of injured striated muscle there are two concurrent processes [23]. The first process is the differentiation of new myofibres from satellite cells. After the injury, activated satellite cells differentiate into myoblasts and fuse with each other into multinucleated myotubes. Myonuclei are located centrally in the myotubes and they are also known as centronucleated cells [20]. The second process of muscle healing is the formation of the granulation and scar tissue, which serve as a scaffold for regenerating myofibres. This process is very similar to the healing of the injured dermis.

Wound tensile strength measurement

Biomechanical studies have shown some clinical and/or experimental importance. Skin sutures can be removed as soon as the wound is fixed enough to no longer need mechanical support (sutures). Therefore, the tensile strength (TS) of wounds is objective and the preferred method for wound healing evaluation is often used in numerous experimental studies [9], [11]. Most of the published papers evaluating the biomechanics have focused only on certain time intervals during the maturation phase of healing [1], [2], [38], [39]. However, the most significant changes occur during the first week of wound healing [34]. In our previous study, we showed that on the body region with minimal strength, there may not be any difference in removing the skin sutures two, three or four days after surgery [13]. Early sutures resection would assure a uniquely better cosmetic result [6].

Histological assessment

It may be suggested that due to the different evaluation techniques and models used, similar works have sometimes resulted in different outcomes. Hence, the use of a standardized and reproducible model is preferable in order to obtain objective information of the experiments. Numerous studies have been done only in a descriptive manner, making a qualitative assessment of the effects of various external factors on the repair processes [11], [40] difficult to evaluate. However, by using this method it is not possible to precisely evaluate the range of changes induced by the tested therapies. Therefore, for the objective evaluation of the effects of various external factors on wound healing, semi-quantitative or quantitative assessments are needed allowing statistical comparison between experimentally treated and control wounds.

It has been shown that there are only quantitative differences between primary sutured and open wound healing. Therefore, for the semi-quantitative histological evaluation of incisional wound healing, a 4-point scale [52] was used, whereas the excisional wound healing was scored by using a 5-point scale [14]. Such scale expansion allows more precise quantification of the occurred changes.

Fluorescence spectroscopy and wound healing

Fluorescence spectroscopy is a type of spectroscopy which analyzes fluorescence from a sample. It involves using a beam of light of certain wavelength and energy that excites the electrons in the studied molecules and causes them to emit light of a longer wavelength and lower energy [45].

The parameters that reflect blood circulation and oxygen delivery in healing tissues have been considered to be the most important variables in wound healing stratification [21]. It is well documented that an increase of oxygen tension reflects an enhanced flavin adenine dinucleotide (FAD) and decreases the reduced nico-

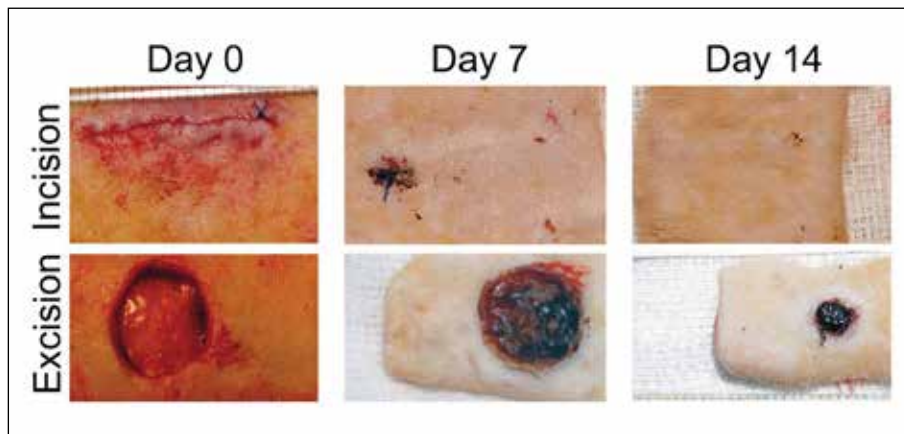


Fig. 1. Basic *in vivo* models of skin wound healing

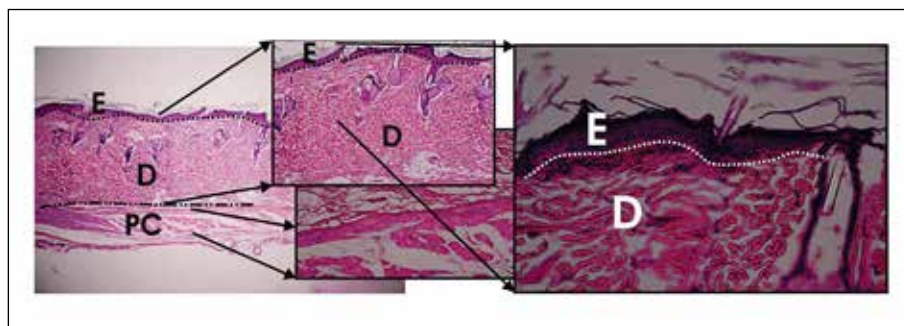


Fig. 2. Histology of rat skin. E — epidermis; D — dermis; PC — *panniculus carnosus*;

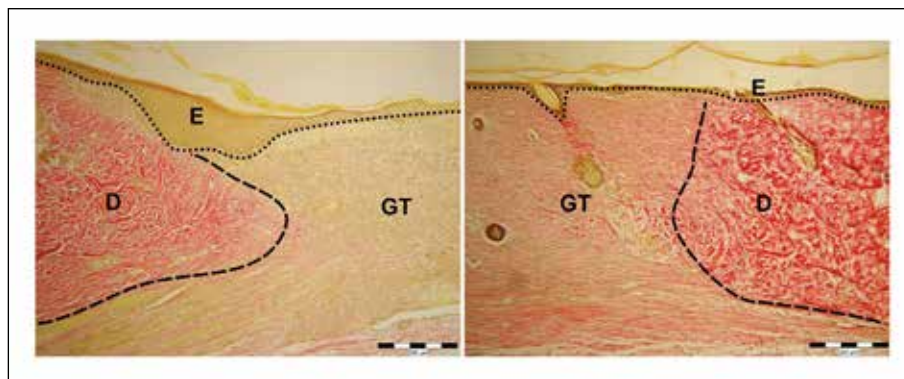


Fig. 3. Skin wounds at one week after surgery

Left — control untreated wound with poor presence of collagen. Right — laser treated wound with granulation tissue rich on new collagen
(E — epidermis; D — dermis; GT — granulation tissue)

tinamide adenine dinucleotide (NADH) concentration; whereas an inadequate oxygen supply elicits the opposite changes in these cofactors [31]. Similarly, increasing the intensity of anaerobic metabolism and lactate production in oxygen insufficiency also leads to a decrease of interstitial pH. Diagnostic techniques based on optical spectroscopy have the potential to monitor biochemical, metabolic, and morphological changes of different tissues. The main advantages of those methods, such as non-invasivity and speed of measurement, have been confirmed in numerous studies [42].

Our spectrofluorimetric study for the first time presented a non-invasive model to investigate the dynamics of selected redox parameters during the early phases of skin wound healing using fluorescence spectroscopy [32]. However, the basic limitation of this study was that it was possible to measure only relative changes and not absolute values of selected redox parameters such as pH, FAD, and NADH. Particularly, the fluorescence spectra in FAD and NADH bands have been shown to be an effective way of monitoring the skin flap ischemic area visualizations [33].

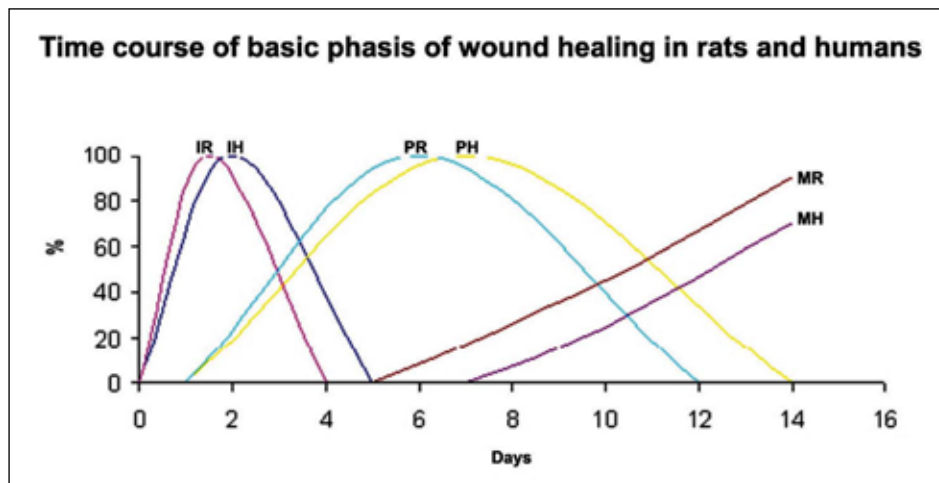


Fig. 4. Comparison of the time courses of basic skin wound healing phases in humans and rats

IR — inflammation rat; IH — inflammation human; PR — proliferation rat; PH — proliferation human; MR — maturation rat; MH — maturation human

Low-level laser therapy (LLLT) and wound healing

Laser (Light Amplification by Stimulated Emission of Radiation) is a device which emits electromagnetic radiation through a process called stimulated emission [17]. Laser light is usually coherent and monochromatic. Low-level laser therapy (LLLT) is a medical technique in which exposure to laser light might modulate cellular function leading to beneficial clinical effects. Rationales for suggesting particular combinations of wavelength, power density, and dose for the treatment of different conditions are still being discussed [7]. Nevertheless, the exact mechanism of action is still unknown, but it is generally accepted that the mechanism is photobiochemical rather than heat-related. An *in vitro* experiment [24] has shown that all tested laser light wavelengths, such as 633, 670, and 820 nm, similarly oxidized cytochrome c-oxidase in HeLa cells. Accordingly, it may be suggested that the molecular mechanism of LLLT action is based on the increase of oxidative metabolism in mitochondria [24].

The LLLT experiments varied in the parameters used for laser radiation, wound models, and treatment protocol [5], [11], [19], [26], [30]. For instance, the assessment of the effect of GaAlAs laser of 904 nm wavelength at 33 J.cm⁻² on days 3, 7, and 14 using an excisional model showed that LLLT accelerated the inflammatory process, epithelization, and positively influenced the collagen deposition in the wounds of both steroid treated and steroid untreated rats [40]. Similarly, increased wound tensile strength following LLLT at 830 nm and 5 J.cm⁻² in diabetic mice was shown [47]. However, in this experiment only two specific days of the maturation phase of healing (11th and 23rd day) were evaluated. In our previous studies [12], [13] we found an accelerated process of both inflammation and proliferation which is in agreement with previously published morphological studies [5], [34]. Moreover, in LLLT (670 nm, dose 5 J.cm⁻²) treated wounds, an increase in the amount of collagen (Fig. 3) and improved epithelization were shown [27] and these results corresponded with the results of experiments where laser radiation of an InGaAlP (685 nm, dose 2.5 J.cm⁻²) laser was used for the healing improvement [46]. The beneficial effects of treatment

with the 980 nm GaAlAs diode laser at 18 J.cm⁻² on wound healing were demonstrated in diabetic mice [25]. On the other hand, treatment at 36 J.cm⁻² had rather adverse effects and decreased the healing rate.

In contrast to those results, in numerous studies, evidence has been provided on the inefficiency of LLLT [37], [41], [44]. In an experimental *in vivo* study, trying to positively affect the healing of burns in rats after using laser irradiation at 635 and 690 nm (1.5 J.cm⁻²), no macroscopic and microscopic differences among treated and control groups were observed [44]. Similarly, no significant differences in epithelization and contraction of skin wounds in horses after LLLT using a GaAlAs laser at 2 J.cm⁻² were demonstrated [41]. In addition, no significantly beneficial effects on the blood microcirculation in wound healing in the laser treated (HeNe laser at 1.5 J.cm⁻²) groups in comparison with controls were demonstrated.

One of the reasons for the negative effect of LLLT in numerous studies may be related to the use of extremely low doses [49]. On the other hand, it has also been shown that power densities above 20 mW.cm⁻² temporarily inhibit fibroblasts metabolism [50]. However, results from our previous study showed that a dose of 30 J.cm⁻² (670 nm) and power density of 25 mW.cm⁻² would be capable of reducing inflammation without compromising fibroblast metabolism [13]. The high variability of the presented results from different studies indicates the need of further detailed assessment of LLLT parameters (wavelength, power density and dose) to achieve the best possible results in promoting the healing process in patients.

DISCUSSION

The relationships between low-level laser therapy and its effect on different models of skin wound healing are still poorly understood. Nevertheless, it has been well demonstrated that both red and infra-red lasers are able to improve wound healing using selected parameters.

Previously, an inverse relationship between wavelength and intensity in red lasers (670 nm vs. 685 nm) has been described at the histological level [11]. In this study, LLLT was found more effective when combining higher intensity with shorter wavelengths or lower intensity with longer wavelengths. The suggestion regarding the inverse relationship of LLLT parameters was partially supported in our biomechanical study (635 nm vs. 670 nm) by using an incisional sutured wound model [51]. However, our observation at 635 nm was observed in their study at 670 nm [11] while their observation at 685 nm [11] was recorded in our study at 670 nm. In addition, in our studies, realized with the excisional model, it was shown that LLLT at both tested wavelengths (635 nm vs. 670 nm) demonstrated similar power density dependent effects [16], [27].

The main difference between incisional sutured and excisional wound healing is in the amount of granulation tissue that is formed during the proliferative phase of healing. Whereas a primary wound heals either without granulation tissue or with a minimal amount of granulation tissue, a secondary wound needs new tissue development which includes extensive granulation tissue formation [13], [14]. Accordingly, we suggested in our previous study that an incisional sutured skin wound should be treated differently than an open wound [36]. In this context, it may be suggested that modern therapy should be wound-type specific.

CONCLUSION

In conclusion, the results from previously published studies showed that the healing of rat epidermis, dermis and striated muscle is, faster than but comparable to, the healing of human skin (Fig. 4) and striated muscle. Hence, rats may be considered as a fair model for basic research of skin repair and its common use by many other authors is well-founded.

The results summarized in this review may be found helpful for a better understanding of selected processes occurring during skin wound healing. In addition, results published by our research group in the field of optical diagnostics and low-level laser therapy should have the potential to improve medical technologies and finally patients' care in veterinary and human medicine. Based on previously observed differences during the treatment of excisions and incisions, it may be suggested that wound therapy should differ for each type of wound.

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EFFECT OF *AESCULUS HIPPOCASTANUM* EXTRACT AND AESCIN ON SOME REACTIVE NITROGEN SPECIES

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ABSTRACT

Reactive nitrogen oxide species, such as highly reactive hydroxyl, peroxy or hydroperoxy radicals, peroxynitrite or hypochlorite, are able to initiate the chain reaction of lipid peroxidation and cause a spectrum of human diseases. Some plant-derived chemicals exhibit very good antioxidant properties, mostly attributed to their high polyphenolic content. The activity of such investigated compounds is attributed mainly to the induction of nitric oxide production, a diffusible free radical. We investigated the ability of dry horse chestnut extract (*Aesculus hippocastanum*) and its main component aescin (escin) at different concentrations to counteract nitric oxide oxidation products and to scavenge peroxynitrite radicals *in vitro*. Aescin showed a better ability to entrap nitrites than the dry horse chestnut extract. Both extracts showed very low effect towards peroxynitrite. At lower concentrations, they even induced peroxynitrite formation. This knowledge may help explain the side effects, but also the persistence of the disease, despite treatment.

Key words: aescin (escin); horse chestnut; nitric oxide; peroxynitrite

INTRODUCTION

“The biology of nitric oxide (NO) has received enormous attention over the past decade due to the ability of this diatomic

molecule to mediate a vast number of physiological processes, such as regulation of vascular tone and neuronal function” [9]. However, at high concentrations of NO (micromolar), the production of reactive nitrogen oxide species (RNOS) becomes significant. Nitric oxide reacts rapidly with superoxide radicals ($O_2^{\cdot-}$) to form peroxynitrite (ONO_2^-), which is, in itself cytotoxic and, readily decomposes into the highly reactive and toxic hydroxyl radical ($\cdot OH$) and nitrogen dioxide (NO_2) [9]. ONO_2^- is much more reactive than NO or $O_2^{\cdot-}$, which causes diverse chemical reactions in biological systems, including nitrosylation reactions which can modify protein structure and inhibit their normal function [6]. Venotonic drugs (horse chestnut and aescin) have been used in chronic venous insufficiency and also in the treatment of diabetic retinopathy [4]. Aescin, the main active compound in horse chestnut seeds, is responsible for most of its medicinal properties. Its effects could be due to the well-known ability of aescin to enhance cellular permeability to calcium which, on one hand, would increase endothelial nitric oxide synthase (eNOS) activity and NO production, and on the other hand, would lead to vascular smooth muscle contraction. Endothelial protection could contribute to the therapeutic usefulness of aescin, while arterial vasoconstriction could represent a limiting side effect in susceptible patients [3]. However, even a short-term increase in NO can lead to chronic illness, presumably by initiating a biochemical vicious cycle recently called the NO/ ONO_2^- cycle [8]. Therefore, an arterial vasoconstriction is not the only possible side effect that could be expected. Moreover, aescin acted as an inhibitor of cell proliferation and an inducer of apoptosis in carcinoma cells [10].

Due to the fact that NO affects several regulation processes even at nanomole concentrations and the resultant substances are capable of modulating the NO production, their overproduction can lead to negative consequences from nitrosative and oxidative stress conditions in biological systems. Thus, the aim of our study was to evaluate the basic ability of the dry chestnut extract and aescin to convert NO to nitrite and scavenge or release ONO_2^- in a dose dependent manner *in vitro*.

MATERIALS AND METHODS

The horse chestnut (*Aesculus hippocastanum* L.) dry extract (HCE, 18 to 22 % aescin) and the saponin beta-aescin (E) were a gift from CALENDULA a.s. (Slovak Republic). The ability to convert NO was estimated indirectly by using the Griess reaction [7], which involves the spectrophotometric measurement of nitric oxide stable decomposition products, i.e. the nitrites and nitrates. NO can be readily oxidized to NO_2 , so that the basic Griess reaction is used as an indirect assay for NO. NO_2 can be further oxidized to NO_3 which does not form the azo dye, but the kinetics of NO_2 oxidation is relatively slow and is therefore, considered to be insignificant [5]. In the aqueous phase, free of biological material, NO exclusively autooxidizes to nitrite [11], so that only the nitrite determination was performed without the need to reduce nitrates prior to the assay. In the Griess reaction, nitrite reacts with sulphanilic acid under acidic conditions to form a diazonium ion which couples to N-(1-naphthyl) ethylenediamine dihydrochloride (NEDD) to form a readily water-soluble azo dye which is detected at 546 nm. The reaction mixture, containing sodium nitroprusside (10 mmol.l^{-1} , 2 ml), phosphate buffer solution (PBS, 1 ml, pH 7.4), and the extracts of different concentrations ($5, 25, 50, 75, 100 \mu\text{g.ml}^{-1}$) was incubated at 25°C for 150 min in the dark. After that, the sulphanilic acid reagent (0.1 % in 5 % H_3PO_4) was added and the mixture was allowed to stand for 8 min, to complete the diazotization. Then, 0.3 ml of NEDD (0.01 %) was added, mixed and allowed to stand for another 8 min in the darkness at 25°C , until a pink colored chromophore was formed.

Peroxynitrite (ONO_2^-) was prepared by mixing 1 volume of cooled H_2O_2 (0.7 mol.l^{-1}) in HCl (0.6 mol.l^{-1}) with 1 volume of cooled potassium nitrite (0.6 mol.l^{-1}), and then 1 volume of cooled NaOH (1.2 mol.l^{-1}) was added to the mixture, as prescribed by Beckman et al. [1]. The reaction mixture was then left to stand overnight under -20°C . The ONO_2^- yield was incubated with the tested plant extracts (diluted in PBS with pH 7.4) in final concentrations, as mentioned above, for 15 minutes and the absorbance of the sample was recorded at 302 nm.

All experiments were performed in triplicate. The percentage inhibition of the reactive nitrogen species generation was evaluated by comparing the absorbance values of the control and experimental tubes and was calculated according to the following equation:

$$(\%) = [(A_{\text{control}} - A_{\text{sample}}) / A_{\text{control}}] \times 100$$

RESULTS AND DISCUSSION

NO has been implicated in many important physiological roles, such as: cytotoxic mediation of the immune system;

regulation of vasomotor tone in the cardiovascular system; and as a neurotransmitter in the central nervous system [12]. The effects of the compounds, particularly of aescin, are related to the production of NO through the enhancement of the cellular permeability to calcium, which would increase eNOS activity [3]. The actual induction and increase in nitric oxide synthase (NOS) activity in the body under certain conditions may not only lead to increased production of NO, but also to the formation of O_2^- [8], which rapidly reacts with NO to form ONO_2^- and thus create conditions leading to the expression of adverse effects, as well as disease chronicity. It is, therefore, important to know the basic effects of the extracts against these forms of reactive species, which would help to explain the mechanisms supporting the clinical indications and also, the possible suppression of undesirable symptoms.

Activities of compounds connected with NO can be assessed indirectly by measuring the NO oxidation products by the Griess reaction. Nitrite and nitrate are stable metabolites of NO, with nitrate being by far the major oxidative metabolites of NO. Because nitrate itself is biologically completely inactive, in contrast to nitrite, it was more convenient to monitor the behaviour of extracts in the aqueous phase free of biological material, where NO exclusively autooxidizes to nitrite [11].

The incubation of a solution of sodium nitroprusside in a phosphate buffer resulted in a linear concentration dependent upon nitrites production, which were reduced by the tested substances (Fig. 1). Both substances counteracted NO oxidation end products, although aescin showed a higher scavenging capacity. The suppression of the released NO may be partially attributed to direct NO trapping, as the tested substances decreased the amount of nitrite generated by the decomposition of sodium nitroprusside *in vitro*. The percentage of NO trapped by aescin increased from $5\text{--}50 \mu\text{g.ml}^{-1}$ (32 %, 37 % and 41 %, respectively) whereas higher concentrations decreased its ability. Horse chestnut showed about twice lower ability to counteract NO end products when compared to aescin.

Although NO is a free radical, it is remarkably nonreactive toward biomolecules. The reaction of NO with O_2 or O_2^- generates RNOS (especially the oxidant product ONO_2^-) which are highly labile and capable of modifying a wider range of biomolecules than NO itself [9]. Both extracts had no significant effect on ONO_2^- scavenging (Fig. 2). While HCE showed very low non concentration dependence on ONO_2^- scavenging (3 %); on the contrary, aescin potentiated the release of ONO_2^- at concentration lower than $25 \mu\text{g.ml}^{-1}$ (Fig. 2). In the case of the whole horse chestnut extract, this may not be a serious finding. It was shown that even the lowest concentration of *A. hippocastanum* extract ($20 \mu\text{g.ml}^{-1}$) is still capable of counteracting O_2^- and OH and thus preventing oxidative and nitrosative stress [2]. The ONO_2^- formation is thus actually hindered by the O_2^- uptake. However, in the case of aescin, the concentration-dependent increase in ONO_2^- is a significant finding. Most importantly, because ONO_2^- is active in oxidizing tetrahydrobiopterin, a cofactor of NOS, NOS deprived of cofactors becomes uncoupled, producing

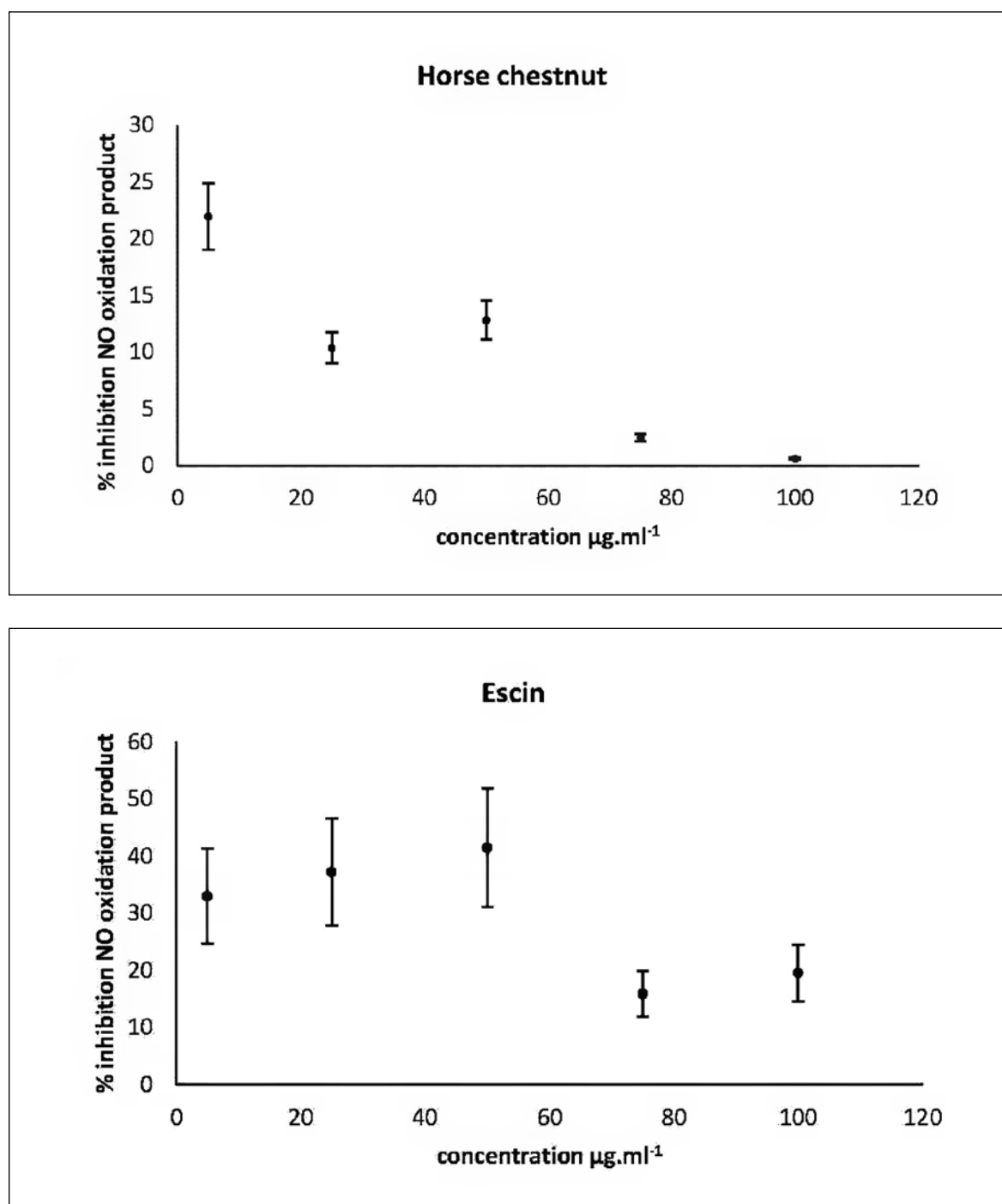


Fig. 1. Effect of horse chestnut extract and aescin (escin) on nitrite, the nitric oxide oxidation product

$\text{O}_2^{\cdot-}$ in place of NO [8]. The critical NO concentration is cell and tissue specific as well as process dependent. The observed conditions of ONO_2^- formation after the induction of NOS by aescin in biological systems indicate that instead of treatment, this would lead to disease chronicity as a result of inadequate dosages. Moreover, the production of ONO_2^- can have pathological consequences because of strong oxidative nitrosylation of molecular targets.

CONCLUSIONS

According to the present knowledge about the effects of extracts of horse chestnut *in vivo*, and the low ability of ONO_2^- uptake, as detected *in vitro*, the low ability to even induce ONO_2^- (in the case of aescin), may form the basis of certain types of therapeutic efficiency; especially when cytotoxic effects against tumour cells are requested, but also of documented side effects at the treatment of venous insufficiency.

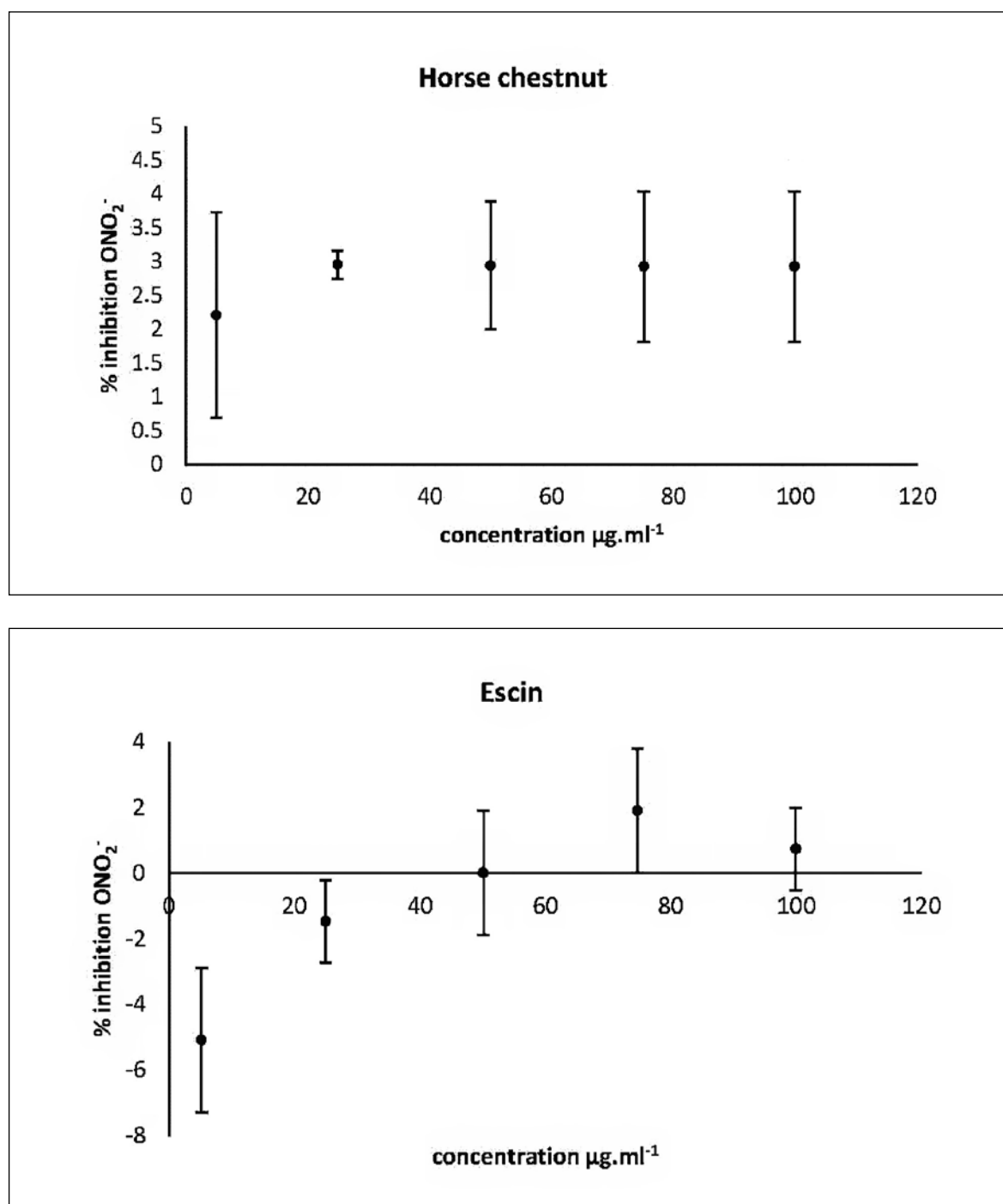


Fig. 2. Effect of horse chestnut extract and aescin (escin) on the exogenous peroxynitrite

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A REVIEW OF GLUTATHIONE PEROXIDASE ROLE IN THE ANTIOXIDANT PROTECTION OF THE ANIMAL ORGANISM

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ABSTRACT

The selenoproteins are the most efficient endogenous components of the antioxidant defence system against reactive oxygen metabolites and their toxic products. Glutathione peroxidase (EC 1.11.1.9, GPx) was the first identified and is the most abundant selenoprotein in mammals. This review considers its structure, distribution, biological function, and the effects of selenium and vitamin E concentrations, on the activity of GPx. Four glutathione peroxidase enzymes have been found in different cell fractions and tissues of the body (cytosolic glutathione peroxidase, GPx-1; gastrointestinal glutathione peroxidase, GPx-2; plasma glutathione peroxidase, GPx-3; and phospholipid hydroperoxide glutathione peroxidase, and GPx-4). Epididymal secretory glutathione peroxidase (GPx-5) and Se-independent GPx occupy a special position in the GPx family. GPx-5 is a secreted enzyme with the absence of the selenocysteine residue at the active site. Selenium-independent GPx is the glutathione S-transferase which can catalyze the reduction of organic hydroperoxides, but not hydrogen peroxide. The functions of the GPxs were originally studied by changing dietary selenium levels. Selenium from organic sources is more efficiently incorporated into tissues than those from inorganic ones. The body's selenium status is assessed by the direct measurement of selenium concentration in blood and tissues, or indirectly by measuring the activity of GPx. The nutrition is one of the important factors which, besides selenium, influence enzyme activity. The effects of selenium deficiency in farm animals, which has been linked to

many health problems mainly in young animals, also have been known to reduce the productive and reproductive abilities of these animals. The deficiency of selenium may also result in immune and endocrine disorders, especially thyroid dysfunction.

Key words: antioxidants; glutathione peroxidase; selenium; selenoproteins; vitamin E

INTRODUCTION

Reactive oxygen and nitrogen species are constantly produced during normal aerobic metabolism. The high reactivity and limited period of existence of free radicals (e.g. superoxide, nitric oxide and hydroxyl) and other reactive species (e.g. hydrogen peroxide, hypochlorous acid and peroxynitrite) play a major role in physiological and pathological processes. The imbalance between the production of reactive oxygen species and the defensive ability of biological systems to detoxify these reactive intermediates causes oxidative stress that can be an important mediator of damage to essential components of the cell (proteins, phospholipids of cell membranes, and DNA). Normally, the body is protected against reactive oxygen metabolites and their toxic products by a wide range of known defensive antioxidant mechanisms, such as: 1. prevention of radicals' formation (metal-binding proteins, superoxide dismutase, glutathione peroxidase, and catalase); 2. prevention and restriction of chain reaction formation and propagation of free radical reactions (vitamin A, E, C, carotenoids, ubiquinol, gluta-

thione, and uric acid); and 3. excision and repair of damaged parts of molecules (lipases, proteases, peptidases, nucleases, and transferases). Oxidative stress has been implicated in the pathology of several diseases, such as; nutritional muscular dystrophy, exudative diathesis, pancreas atrophy, hepatitis dietetica, etc. Therefore, oxidative stress must be controlled by supplying all known antioxidant nutrients and by minimizing the effects of substances that stimulate reactive oxygen species [25].

THE SELENOPROTEINS

An organism defends itself against the negative effects of free radicals by different antioxidant mechanisms and mainly the selenoproteins represent the most efficient endogenous component of this defence system. It has been suggested that up to 100 selenoproteins may exist in mammalian systems, of which up to 30 have been identified by ^{75}Se labelling *in vivo*. To date 15 selenoproteins have been purified, or cloned, allowing further characterization of their biological functions. The biological function of selenium in mammals has been studied mainly in the context of its antioxidant role in protecting tissues from oxidative degeneration. Selenium, an essential trace element, is present in all cells and tissues as a component of several major metabolic pathways, including; reproduction and thyroid hormone metabolism, antioxidant defence systems, and immune functions [43].

The role of selenocysteine

Biosynthesis of all the selenoproteins depends on the availability of selenium which is incorporated as selenocysteine (Sec), the 21st amino acid at the active site of a wide range of selenoproteins. The importance of selenocysteine comes from its role in gene expression and the enzyme activity of the Se-dependent GPx [19]. Selenocysteine is synthesized from the activated selenium, selenophosphate and activated serine residue (Ser) on Sec^{tRNA} (Ser Sec^{tRNA}) to Sec Sec^{tRNA} . The codon for the incorporation of Sec residue is UGA that is the stop codon in general. Five components are known to be required for the translation of UGA codon to Sec residue in selenoproteins: two *cis*-sequences (SECIS — a Sec insertion sequence element in the 3'-untranslated region, and a Sec codon, UGA, in the coding region) and three known *trans*-acting factors (EPSeC — a Sec specific translation elongation factor, the Sec Sec^{tRNA} , and SBP2 — a SECIS binding protein) [28].

All of the selenoproteins for which an enzymatic activity has been identified catalyze redox reactions involving oxidation of thiol groups and/or reduction of disulphides. Selenium is isomorphous with sulphur, having the same outer electron structure, a slightly larger atomic radius and essentially identical electronegativity. Thus, selenium homologues exist for most sulphur-containing molecules. Despite these structural similarities, selenium is a much stronger nucleophile than sulphur, its compounds have much more negative redox potentials than homologous sulphur compounds, and hydrogens bonded to selenium are much more acidic than those bonded to sulphur ($\text{pK}_a \sim 5$ vs. ~ 9). Because of these properties, selenocysteine is able to catalyze the oxidation of thiols in the reducing environment of the cytoplasm, where the reduced glutathione (GSH) to oxidized glutathione (GSSG) ratio of 30–100 to 1 results in a redox potential of approximately -230 mV [47]. Sele-

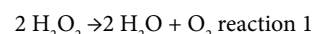
noproteins have the unique ability to trap H_2O_2 and use the resulting selenenic acid to form specific disulfide bonds in the presence of $10\text{--}20\text{ mmol.l}^{-1}$ GSH and are thus well suited for controlling the formation of disulfide bonds with specific Cys residues. Because of the rapid oxidation kinetics of selenols, H_2O_2 in the cytosol reacts completely with Sec-containing proteins to produce Sec selenenic acid before it can react with other potential targets [46]. Thus, rapid consumption of H_2O_2 by selenoproteins acts to limit the duration of H_2O_2 signals and the distance over which those signals are transmitted by diffusion.

GLUTATHIONE PEROXIDASE

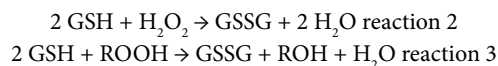
Glutathione peroxidase (EC 1.11.1.9, GPx) is the first identified and the most abundant selenoprotein in mammals. The first report of this glutathione peroxidase activity was made by Mills [31] who demonstrated glutathione dependent catabolism of hydrogen peroxide by a bovine red blood cell lysate. In erythrocytes, GPx removes H_2O_2 formed by dissociation of oxyhaemoglobin into O_2^- and methaemoglobin [32]. The enzymology and the role of glutathione peroxidase was first described by Flohé and Schlegel [9], while Rotruck et al. [40] isolated this enzyme from rat liver and confirmed that it contained selenium. The major physiologic role of this selenoenzyme is to protect biomembranes and other essential cellular components against oxidative challenge and to maintain appropriately low levels of hydroperoxides within the cell, thus decreasing potential free radical damage.

Catalytic mechanism for reduction of hydroperoxides by glutathione peroxidase

In an aerobic organism, hydrogen peroxide is catabolized by different enzyme systems. High H_2O_2 concentrations are scavenged by catalase (EC 1.11.1.6, CAT) which catalyzes very efficiently dismutation of hydrogen peroxide to form water and molecular oxygen (reaction 1).



At low concentrations of H_2O_2 , glutathione peroxidase catalyzes its reduction using glutathione (reaction 2) and this enzyme is also capable of reducing a variety of hydroperoxides including lipid hydroperoxides (reaction 3) to water and the corresponding alcohols at the expense of the reducing equivalent glutathione, which serves as a specific electron donor substrate [10].



The GPx catalytic site includes a selenocysteine residue in which the selenium undergoes a redox cycle involving the selenol (ESeH) as the active form that reduces hydrogen peroxides and organic peroxides (Fig. 1). The selenol is oxidized to a selenenic acid derivative (ESeOH), which reacts with reduced glutathione (GSH) to form a selenenyl sulphide complex (ESeSG) consisting of the enzyme-bound selenium and GSH. A second glutathione molecule then reduces this intermediate back to the selenol (ESeH) releasing the oxidized glutathione (GSSG) as the by-product [41]. The

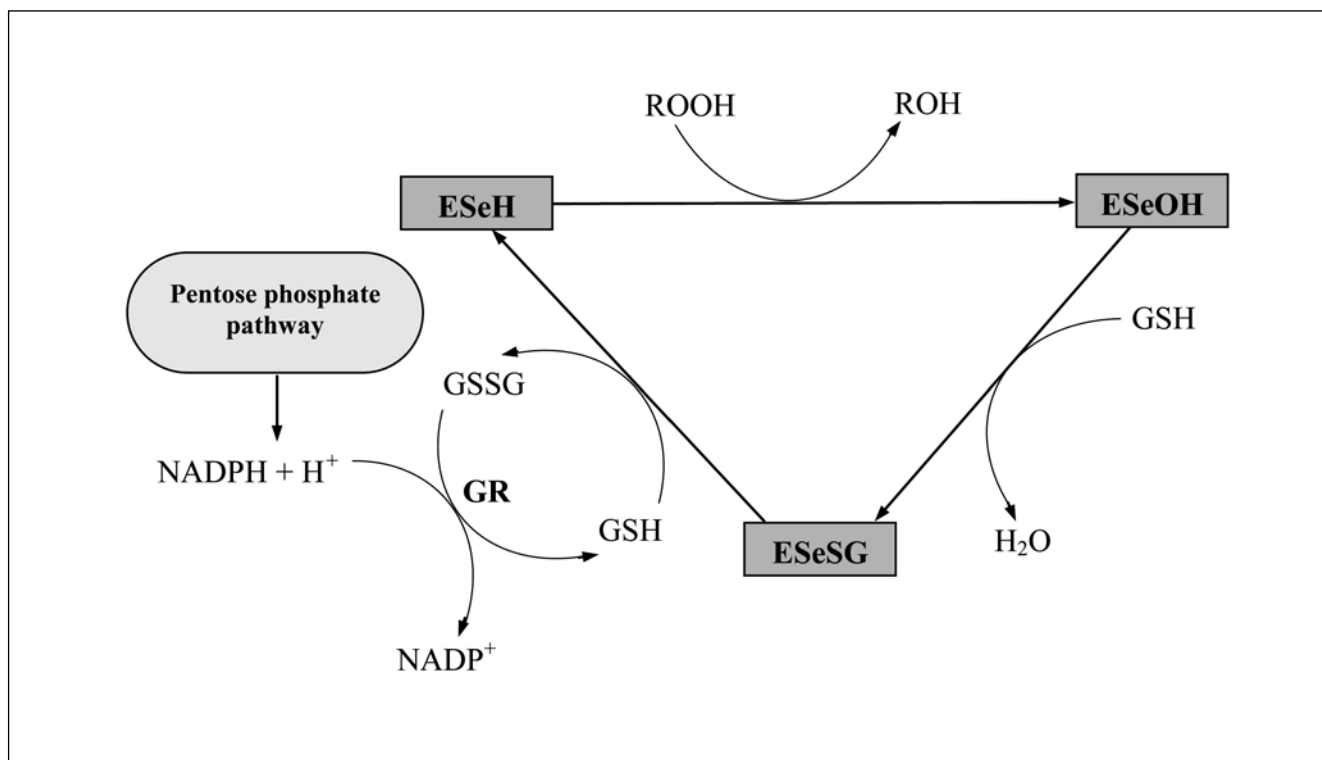
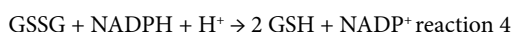


Fig. 1. Catalytic mechanism for reduction of hydroperoxides by glutathione peroxidase
(ESeH — selenol; ESeOH — selenenic acid; ESeSG — selenyl sulphide complex;
GSH — reduced glutathione; GSSG — oxidized glutathione; GR — glutathione reductase)

oxidized glutathione is reduced by the flavin adenine dinucleotide-dependent enzyme, glutathione reductase (EC 1.8.1.7, GR), to the thiol form GSH to complete the cycle. NADPH+H⁺ reduces FAD present in GR to produce a transient FADH⁻ anion quickly breaking a disulfide bond (reaction 4). The oxidation of NADPH+H⁺ to NADP⁺ is accompanied by a decrease in absorbance at 340 nm [35].



So, for each GSSG, two reduced GSH antioxidant molecules are produced, scavenging reactive oxygen species in the cell. In cells exposed to high levels of oxidative stress, like red blood cells, up to 10% of the glucose consumption may be directed to the pentose phosphate pathway (PPP) for the production of the NADPH+H⁺ needed for this reaction. In erythrocytes, if the PPP does not function, then the oxidative stress will lead to lysis, and eventually anaemia.

Glutathione (GSH, γ -glutamylcysteinylglycine) is the major endogenous antioxidant produced by the cells, participating directly as a free-radical scavenger. The ratio of reduced glutathione to oxidized glutathione within cells is often used as a measure of cellular antioxidant status. When cells are exposed to increased levels of oxidative stress, GSSG will accumulate and the ratio of GSH to GSSG will decrease. Therefore, the determination of the GSH/GSSG ratio and the quantification of GSSG are useful indicators of oxidative stress in cells and tissues [27]. Glutathione is able to regenerate the most important antioxidants, vitamins C and E back to their active forms. The intracellular content of glutathione depends

on environmental factors and functions as a balance between its utilization and synthesis.

THE GLUTATHIONE PEROXIDASE FAMILY

Four structurally and genetically different forms of selenium-containing GPx have been functionally described and exist in different tissues or parts of the cell: cytosolic glutathione peroxidase (cellular, classical, cGPx, GPx-1), gastrointestinal glutathione peroxidase (giGPx, GPx-2), plasma glutathione peroxidase (pGPx, extracellular eGPx, GPx-3) and phospholipid hydroperoxide glutathione peroxidase (phGPx, GPx-4). Although GPx-1, GPx-2, and GPx-3 are homotetramers, GPx-4 is a monomer with a molecular size smaller than the subunits of the other glutathione peroxidases [5]. The small size and its hydrophobic surface have been implicated in its ability to react with complex lipids in membranes.

The role of the individual GPx isoenzymes

All glutathione peroxidases reduce hydrogen peroxide and alkyl hydroperoxides at the expense of glutathione with their different specificities for the hydroperoxide substrate. Whereas cytosolic glutathione peroxidase (GPx-1) reduces hydrogen peroxide and some organic hydroperoxides (cumene hydroperoxide, t-butyl hydroperoxide), phospholipid hydroperoxide glutathione peroxidase (GPx-4) and to some extent plasma glutathione peroxidase (GPx-3) also reduce hydroperoxides of more complex lipids (phos-

phatidylcholine hydroperoxide) [11], [45], [48]. GPx-4, however, efficiently reduces hydroperoxy groups of thymine, lipoproteins, and cholesterol esters and is unique in acting on hydroperoxides integrated in membranes [44].

Gastrointestinal glutathione peroxidase (GPx-2) is the closest homologue of GPx-1. It has approximately 65 % of the amino acid sequence identity and 60 % of the nucleotide sequence identity with GPx-1 [6]. This enzyme represents a major barrier against hydroperoxides originating from nutrition. During feed production and storage, some unsaturated lipids are oxidized and they can cause health-related problems, including decreased growth, impaired reproductive function and immune competence [20]. These oxidized lipids can react with transition metals, already present in the feed or as feed supplements, producing free radicals which can react with natural or synthetic antioxidants present in the feed. GPx-2 prevents them, in this way, entering systemic circulation.

The mRNA for GPx-3 is found predominantly in the kidney, in particular the epithelial cells of the proximal tubules [3] and therefore it shows a significant diagnostic value in kidney diseases. GPx-3 may have a specific antioxidant function in renal tubules or extracellular fluids. In the case of low concentration of glutathione, Björnsted et al. [4] have suggested other thiols such as thioredoxin and glutaredoxin systems which can also act as electron donors and support an antioxidant role for plasma GPx. Investigating selenium deficiency in humans, Cohen et al. [7] found that after selenium supplementation of Se-deficient individuals, plasma GPx-3 activity returned to normal values within 4–5 weeks, whereas 3–4 months were required for the red blood cell enzyme to reach this effect.

Phospholipid glutathione peroxidase decreases lipid hydroperoxide levels, and thus inhibits the lipoxygenases that metabolise arachidonic acid to generate intermediates that mediate signals for increasing cell proliferation and inhibiting apoptosis. In particular, it inhibits 5-lipoxygenase and reduces the production of 5-hydroxyicosatetraenoic acid, which is known to stimulate the proliferation of prostate cancer cells [12]. GPx-4 has an unusual tissue distribution. In the rat testes it is 15 times more active than in the liver and kidneys, and 25 times more active than in the heart and lungs [2]. Recently, Godeas et al. [13] have proposed an additional role for GPx-4 in the maturation of spermatozoa *via* the metabolism of hydroperoxides and sperm thiol oxidation. Phospholipid GSH-Px may also be involved in the regulation of leukotriene biosynthesis.

Epididymal secretory glutathione peroxidase (GPx-5) occupies a special position in the GPx family and it has been demonstrated in rats, mice, pigs, monkeys and humans. The peculiarity of this GPx form is the absence of the selenocysteine residue at the active site. It is replaced by a cysteine amino acid residue. The purified protein contains no selenium, and its GPx activity towards hydrogen peroxide or organic peroxides is less than 0.1 % of that of GPx-1. This low activity, coupled with very low glutathione concentrations in the epididymal fluids, has led to the suggestion that this protein protects sperm by binding organic hydroperoxide which would otherwise induce an acrosome reaction [8].

The glutathione S-transferases (EC 2.5.1.18, GSTs), one of the most abundant liver proteins, play an important role in protection against oxidative stress by mechanisms involving conjugation of aldehydes and other potentially oxidative compounds. This group of enzymes has the primary role in xenobiotic detoxification metabo-

lism by catalyzing the transfer of reduced glutathione to various substrates containing a reactive electrophilic centre, to form a polar S-glutathionylated product for reducing oxidative stress. In addition to conjugation reactions, the glutathione S-transferases can also function as antioxidants to reduce organic hydroperoxides, but not hydrogen peroxide. Therefore, they are sometimes referred to as enzymes with the glutathione peroxidase activity independent of selenium (Se-independent GPxs). Using hydrogen peroxide as substrate, the activity of Se-independent GPx is not measured because this enzyme cannot catalyze the reduction of hydrogen peroxide and thus this activity is only against organic hydroperoxides with a much higher K_m than Se-dependent GPx [16]. Selenium deficiency causes an increase in hepatic glutathione S-transferase activity. In rat liver, glutathione S-transferase is present in the cytosol, mitochondria, and microsomes. Its GPx activity in the microsomes appears to be lower than that in the other sites. Rat liver glutathione S-transferase activity is 50 to 100 % higher in Se-deficient male rat liver cytosol than in the control. Lawrence and Burk [24] reported the presence of both Se-dependent and Se-independent GPx activities in pig livers. Approximately 67 % of the total liver GPx activity of the pig was Se-independent, whereas the data of Meyer et al. [29] suggested that 45 to 60 % was Se-independent. The amount of Se-independent GPx activity was fairly consistent at the different dietary Se levels, the relative percentage of Se-independent activity was greatest for those fed 0.1 ppm Se because of the lower Se-dependent GPx activity. These results suggest that the increase in total liver GPx activity largely reflected the increase of the Se-dependent form of the enzyme, as there was no further increase in activity at higher dietary Se levels.

THE EFFECT OF SELENIUM AND VITAMIN E CONCENTRATION ON THE ACTIVITY OF GLUTATHIONE PEROXIDASE

Selenium is an essential micronutrient element in diets of various animal species, which is required for maintenance of health, growth and biochemical-physiological functions, such as: biological antioxidation, immune function, reproduction and thyroid hormone metabolism. Selenium can be found in all the cells and tissues of the body but its levels in blood and tissues are very much influenced by dietary selenium form and intake. It is also well known that selenium from organic sources is more efficiently incorporated into tissue than inorganic sources of selenium [38]. The body's selenium status is assessed by direct measurement of selenium concentration in blood and tissues, or indirectly by measuring the activity of glutathione peroxidase. Since selenium is incorporated by erythrocyte glutathione peroxidase during erythropoiesis, the glutathione peroxidase enzyme is considered to be a suitable indicator of biologically active selenium and thus an indicator of long-term supplementation of an organism with selenium. The direct determination of the selenium content in the blood reveals the short-term selenium status of the body [37]. These data were confirmed by Kováč and Sankari [23], who found the highest correlation between whole blood selenium concentrations and erythrocyte GPx activities ($r=0.992$), and between selenium concentration and GPx activities in the whole blood ($r=0.981$). Konvičná et al. [21] also observed corresponding GPx activity in the range of 569.32–

691.90 $\mu\text{kat.l}^{-1}$, with the use of the most common reference selenium value (70.30–100.32 $\mu\text{g.l}^{-1}$) in the whole blood of dairy cows, and the correlation analysis showed the degree of dependence between the measured indices ($r=0.892$; $y=0.0031x-0.8749$). In the study of Pavlata et al. [37], the dependence of GPx activity on the selenium concentration in the blood of sheep was found and the activity values of GPx were determined for use in the diagnosis of sufficient selenium status in sheep in the Czech Republic (GPx activity greater than 600 $\mu\text{kat.l}^{-1}$ of whole blood). Ludvikova et al. [26] defined for the practical establishment of selenium deficiency in horses the marginal level of GPx activity as 200–300 $\mu\text{kat.l}^{-1}$ in whole blood, when the deficiency level for selenium is set as 75–100 $\mu\text{g.l}^{-1}$ of Se in the blood. Misurova et al. [34] found that a selenium concentration of 100 $\mu\text{g.l}^{-1}$ corresponds with GPx activity 700 $\mu\text{kat.l}^{-1}$ in the blood of adult goats and 720 $\mu\text{kat.l}^{-1}$ in the blood of newborn kids.

Selenium deficiency causes various diseases; the most frequent are muscle dystrophy in calves, young cattle after turning to pasture, lambs, foals, poultry, hepatosis dietetica in pigs, exudative diathesis, and encephalomalacia in poultry. Moreover, various reproduction disorders occur (infertility, degeneration of seminiferous tubules), reduced libido in sheep, disorders of ovarian cycle, and increased occurrence of infertility [22]. Selenium deficiency has also been linked to many health problems in young animals such as: increased neonatal mortality; decreased sucking reflex; weakness; and higher occurrence of infectious diseases. Selenium, which is essential as a trace element for intrauterine development of calves, passes in considerable amounts through the placental barrier to the foetus, thus reducing the body pool of selenium in a pregnant female. This is why in late pregnancy, when foetuses develop and increase their weights most intensively, cows have much lower serum selenium concentrations compared to non-pregnant heifers and cows [42]. High-producing transition cows are highly susceptible to oxidative stress, which Aitken et al. [1] report to contribute to mastitis pathogenesis by modifying the expression of genes coding for proinflammatory factors. The same authors report that during the dry period, the activity of glutathione peroxidase is relatively low, but increases considerably during early lactation, when it serves a protective role in the proinflammatory state of the mammary gland. As shown by Miranda et al. [33], the reduction in the number of mammary epithelial cells with advancing lactation is at least in part caused by oxidative stress, which is accompanied by the loss of contact and cuboidal structure of cells, and the decrease in their viability as a result of apoptosis stimulation.

Vitamin E, the primary lipid-soluble antioxidant in cell membranes, is part of the body's intracellular defence against the adverse effects of reactive oxygen species and free radicals that initiate oxidation of unsaturated phospholipids and critical thiol groups of proteins. Muscle damage and muscular dystrophy are common signs of both vitamin E and selenium deficiency. The more metabolically active tissues (such as skeletal and smooth muscles and liver) have a greater potential for oxidative tissue damage if the vitamin E supply is limiting. Erythrocytes and capillary walls are also susceptible to damage in animals with marginal vitamin E status. The protection of cell membranes and other cellular components of immune cells against lipid peroxidation is probably the most important mechanism of vitamin E in the immune response. Vitamin E and selenium enhance host defences by improving phagocytic cell function.

Both vitamin E and GPx protect phagocytic cells and surrounding tissues from oxidative attack by free radicals produced by the respiratory burst of neutrophils and macrophages during phagocytosis [36]. Hogan et al. [17] reported that vitamin E supplementation of the diet, increased the intracellular kill of *Staphylococcus aureus* and *Escherichia coli* bacteria by neutrophils. Cows supplemented with 2.01 g vitamin E per day, in the presence of adequate selenium (0.3 ppm) starting four weeks prior to calving through eight weeks postpartum, had increased neutrophil and macrophage functions and reduced somatic cell counts compared to controls [39]. Supplemental vitamin E was shown to specifically stimulate the phagocytosis of *Staphylococcus aureus* by bovine neutrophils. Both vitamin E and selenium increased neutrophil chemotaxis and superoxide production in these studies. Milad et al. [30] suggested that vitamin E and selenium supply led to a significant rise in whole blood GPx activity and differences in phagocytic functions of leukocytes and neutrophils in sheep. Vitamin E also supports the immune function by its effects on arachidonic acid metabolism and subsequent synthesis of prostaglandins, thromboxanes and leukotrienes. The production of these compounds increases under stress conditions. Thromboxane and interleukin-2 appear to exert a negative feedback effect on leukocyte function [14]. In dairy cattle, supplementation with selenium, or both selenium and vitamin E, reduced the incidence of retained placenta in herds where the prevalence of retained placenta was high, or when selenium or vitamin E were marginal in the diet [18]. Supplemental vitamin E-selenium has also been reported to reduce metritis, cystic ovaries and the time of uterine involution in cows with metritis [15]. Zigo et al. [49] observed the effect of parenteral application of selenium and vitamin E on the health of mammary glands in dairy cows during the peripartum period. It is one of the ways of how to prevent their deficit related to supplied feed rations and ensure the adequate concentration of Se and activity of GPx in the blood and thus increase the natural resistance of dairy cows to intramammary infections.

CONCLUSION

Glutathione peroxidase is an important part of the antioxidant defence system and the existence of four Se-dependent GPxs in different cell compartments immediately implies specific functions for each of the isozymes. GPx scavenges most of the endogenously generated H_2O_2 in the cells and so it has a primary physiological importance in protecting the erythrocyte haemoglobin from oxidative breakdown. Since the erythrocyte has a high GSH concentration and adequate mechanisms for maintaining the glutathione in the reduced state, this GPx system provides an explanation for the remarkable stability of haemoglobin in the intact erythrocyte. GPx plays a key role in modulating the GSH/GSSG ratio and indirectly affects the NADP/ NADPH+ H^+ quotient of the cell. The enzyme may thereby regulate multiple cellular functions such as: cell division, pentose phosphate pathway, gluconeogenesis, mitochondrial oxidation of α -oxo-acids and others. Glutathione peroxidase also reduces lipid hydroperoxides, potentially harmful prooxidants that may promote peroxidation of polyunsaturated fatty acids in phospholipids in biological membranes.

The presence of selenium in the active sites of GPx enzyme indicates an important biological role for this trace element in the normal functioning of living cells. This biological function of selenium in mammals has been studied mainly in the context of its antioxidant role in protecting tissue from oxidative degeneration. Selenium enables the functioning of the immune system and prevents development of malignant diseases. Se-dependent GPx inhibited the initiating and promoting stages of chemical carcinogenesis. In some diseases of farm animals, the effects of selenium deficiency reduced their productive and reproductive abilities. Deficiencies of selenium result in biological dysfunction associated with protein or membrane damage. There are several factors abrogating the activity of the enzyme. Some of these are individual internal factors, resulting in a significant variation in the enzyme activity in different organs, body weight, age groups, sex and endocrine regulation. However, environmental factors also have definite effects on enzyme action. Nutrition is one of the most essential factors, as fat content and fatty acid composition of feed, or trace element intake as well as vitamin status of the animal, play a crucial role in normal enzyme activity. Selenium supplementation is required to overcome the deficiency symptoms and the bioavailability of the same depends upon the nature of supplements used. It is generally found that organic selenium compounds have substantially greater bioavailability than that of inorganic selenium and that selenium from organic sources is more efficiently incorporated into tissue than inorganic sources of selenium.

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EFFECT OF HEAVY METALS ON EXPERIMENTAL *TRICHINELLA SPIRALIS* INFECTION IN MICE

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ABSTRACT

Exposure of hosts to heavy metals can weaken their immunocompetence and increase their susceptibility to parasitic infections. Mice were chronically intoxicated with lead (Pb), cadmium (Cd), or mercury (Hg) and subsequently infected with *Trichinella spiralis* larvae. Our aim was to evaluate the effect of heavy metals on the intensity of parasitic infections. Lead did not affect parasite numbers in the intestine, but during the muscle phase of the infection, it did increase the numbers of larvae. Cadmium significantly reduced the intensity of intestinal infections and also the larva infestations of mice. Mercury intoxication prolonged the intestinal phase until day 20 post infection (p. i.), which may be caused by debilitation of the intoxicated animals. The numbers of muscle larvae in the Hg-intoxicated mice, in relation to their weight, were the highest of the heavy metals tested, resulting in the heaviest parasite burden.

Key words: cadmium; heavy metals; lead; mercury; mouse; *Trichinella spiralis*

INTRODUCTION

Parasites have long attracted ecologists' attention because they can serve as potential indicators of environmental quality by determining the risk rate of toxic bio-accumulation in a host. Even though small mammals (due to their high abundance and wide

distribution) are preferably being used as bio-indicators of environmental contamination, their parasites can accumulate contaminants in their bodies in higher concentrations than their hosts. However, very little is known about the effects of their accumulation capacity for heavy metals on the hosts and the effects on the course of parasitic infections in a heavy metal polluted environments and particularly how they affect a host's health.

For the purposes of environmental evaluations, the most suitable entities are the parasites with great geographic range and numerous hosts, such as *Trichinella spiralis* nematodes, who have cosmopolitan distribution in moderate and tropical climate zones [19]. The main hosts of *Trichinella* are domestic pigs (*Sus scrofa domestica*) and wild boars (*Sus scrofa scrofa*), followed by synantropic animals like rats, dogs, cats and certain wild carnivores, such as bears, red foxes and coyotes [4], [3], [20]. Horses are considered unusual hosts, due to their herbivorous diet, however, their infections are highly pathogenic to humans who consume horse meat. *T. spiralis* is the causative agent of trichinellosis, characterized by enteritis (caused by adult worms), inflammation and degenerative changes in the skeletal muscles (caused by the larvae) [9]. The pathology of trichinellosis is essentially a reaction to an initial inflammatory response during the intestinal phase and the subsequent allergic and inflammatory responses during the larval migration and invasion of the host's muscle [17]. Small mammals play a significant role in the transmission of this helminthic zoonosis. In the potentially contaminated environment they present a large host population that becomes an easy prey when weakened by parasitic infection or by heavy metals, thus increasing the risk of infection and concentration of toxic substances in the predators.

Table 1. Experimental scheme

| Group (number of mice) | Intoxication (daily, from day 0 to the end of the experiments) | Infection (per os, on day 30 of experiments) |
|---------------------------|--|---|
| 1 (n = 21) | control | 500 <i>T. Spiralis</i> larvae/mouse |
| 2 (n = 21) | Pb(CH ₃ COO) ₂ ·3H ₂ O 1 mg.l ⁻¹ drinking water <i>ad libitum</i> | 500 <i>T. Spiralis</i> larvae/mouse |
| 3 (n = 21) | Pb(CH ₃ COO) ₂ ·3H ₂ O 100 mg.l ⁻¹ drinking water <i>ad libitum</i> | 500 <i>T. Spiralis</i> larvae/mouse |
| 4 (n = 21) | CdCl ₂ 100 mg.l ⁻¹ drinking water <i>ad libitum</i> | 500 <i>T. Spiralis</i> larvae/mouse |
| 5 (n = 21) | CdCl ₂ 200 mg.l ⁻¹ drinking water <i>ad libitum</i> | 500 <i>T. Spiralis</i> larvae/mouse |
| 6 (n = 21) | HgCl ₂ 0.2 mg.kg ⁻¹ body weight (b. w.), subcutaneously | 500 <i>T. Spiralis</i> larvae/mouse |
| 7 (n = 21) | HgCl ₂ 1.0 mg.kg ⁻¹ b. w., subcutaneously | 500 <i>T. Spiralis</i> larvae/mouse |

The aim of this study was to determine the intensity of *T. spiralis* infections in the host organism intoxicated with lead, cadmium and mercury. The data may elucidate the role of heavy metals in parasitic infections relating to bio-indicative value of its hosts – small mammals in the ecosystems contaminated with heavy metals.

MATERIALS AND METHODS

The experiments were performed on 147 BALB/c male mice, divided to 7 groups with 21 animals in each, weighing 20–22 g. The animals were kept under standard conditions (at 20–21 °C, 50–60 % relative humidity) and fed *ad libitum* using a commercial diet. The mice, except for Group 1, were intoxicated with different doses of heavy metals in the drinking water (Pb, Cd) or subcutaneously with Hg, from day 0 to the end of the experiment and subsequently infected with the same dose of *T. Spiralis* larvae on day 30 of the experiments (Table 1).

Determination of intestinal worm burdens

The intestinal phase of the infection was investigated on days 5, 10, 15 and 20 post infection (p. i.). The small intestine was cut into 5–10 cm long pieces, placed into a sieve and incubated in conical Pilsner glasses in 0.9 % saline at 37 °C overnight. After incubation, the gut pieces were discarded and the sediment was examined under a stereomicroscope at 60 x magnification.

Isolation of muscle larvae

The muscle phase of infection was examined on days 20 and 30 p. i. Whole eviscerated carcasses were minced and artificially digested (1 % pepsin, 1 % HCl for 4 h at 37 °C; both Sigma-Aldrich, Germany) according to Kapel and Gamble [10]. The samples were allowed to settle for 20 min before the supernatant was discarded and the sediment was poured through a 180 µm sieve into a conical glass and washed with tap water. The sediment was finally transferred to a gridded Petri dish and counted using a stereomicroscope at 40x magnification. Depending on the density of larvae, either a sub-sample or the whole sample was counted.

Statistic evaluation was carried out by one-way ANOVA method and the differences were considered significant at $P < 0.05$.

RESULTS AND DISCUSSION

The animals were intoxicated with different doses of heavy metals throughout the whole experiment (60 days) and infected with *T. spiralis* after being intoxicated for 30 days. During the intestinal phase, the adults were isolated from the small intestine on days 5, 10 and 15 p. i.; no parasites were recovered from the small intestine on day 20 p. i. in all groups except for the groups intoxicated with mercury (Fig. 1). The group with the higher mercury exposure (1 mg.kg⁻¹ b. w.) displayed the highest increase in the number of parasites in the small intestine on day 10 p. i., compared to Group 1. The groups with Hg exposure had the highest number of parasites on days 5 and 10 p. i. The mercury-exposed and *T. Spiralis* infected mice suffered from cachexia, which may have affected the host parasite load.

Compared to non-intoxicated mice, the exposure to lead did not affect the intensity of the parasitic infection. On day 15 p. i., mice intoxicated with 100 mg.l⁻¹ of lead acetate had lower numbers of parasites than the non-intoxicated mice, though the differences were insignificant.

The parasite load was reduced in mice exposed to cadmium and a number of parasites in the small intestine in these mice was significantly lower ($P < 0.05$) at the lower cadmium dose of 100 mg.l⁻¹ on days 5 and 10 p. i.

No significant differences in larva burden were observed when assessing the muscle phase of the infections (Fig. 2). The muscle larvae were found on day 20 p. i.; on day 30 p. i. a higher number of larvae were found in the groups subjected to lead and cadmium intoxication at a dose of 100 mg.l⁻¹. A reduced number of adults in the mice intoxicated with cadmium (200 mg.l⁻¹) resulted in a lower number of muscle larvae, though the differences were statistically insignificant. Mercury intoxication on day 20 p. i. resulted in a very

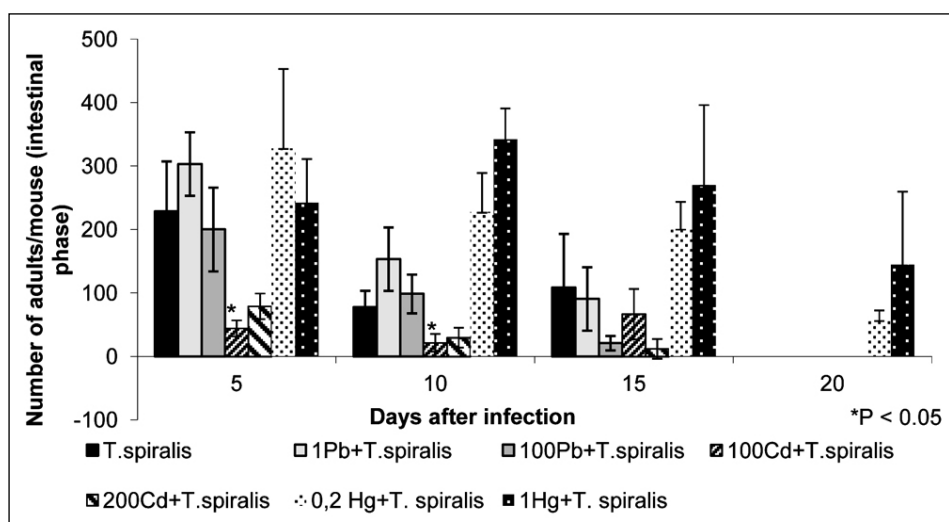


Fig. 1. The intensity of parasitic infection in the intestinal phase of trichinellosis in mice intoxicated with heavy metals (Pb, Cd, Hg) on days 5, 10, 15 and 20

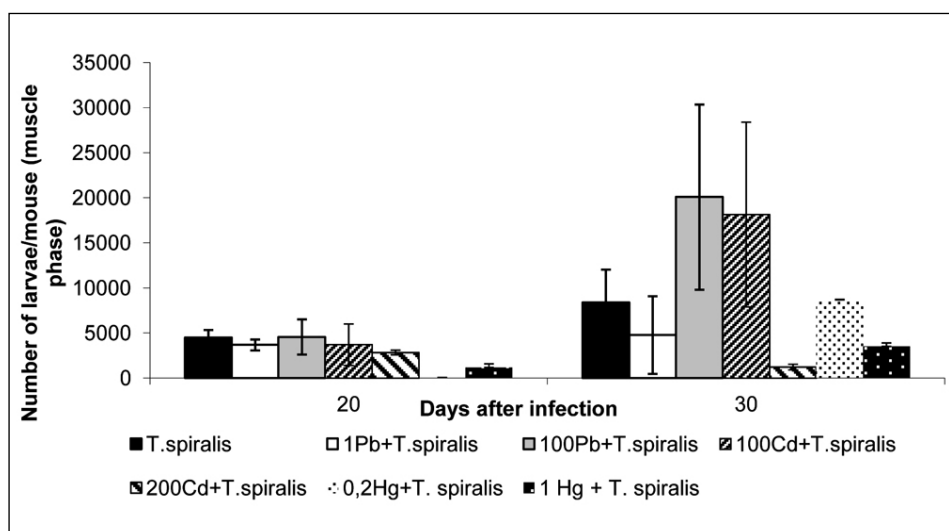


Fig. 2. The intensity of parasitic infection in the muscle phase of trichinellosis in mice intoxicated with heavy metals (Pb, Cd, Hg) on days 20 and 30

Table 2. Index of reproductive capacity (RCI) in mice intoxicated with heavy metals (Pb, Cd, Hg) and infected with *T. spiralis*

| | G-1 | G-2 | G-3 | G-4 | G-5 | G-6 | G-7 |
|-----------|--------------------|---------------------------------|-----------------------------------|-----------------------------------|-----------------------------------|-----------------------------------|-----------------------------------|
| Day p. i. | <i>T. spiralis</i> | 1 Pb + <i>T. spiralis</i> | 100 Pb + <i>T. spiralis</i> | 100 Cd + <i>T. spiralis</i> | 200 Cd + <i>T. spiralis</i> | 0.2 Hg + <i>T. spiralis</i> | 1.0 Hg + <i>T. spiralis</i> |
| 20 | 9.00 | 7.37 | 9.13 | 7.43 | 5.70 | 0.08 | 2.39 |
| 30 | 16.80 | 9.57 | 40.17 | 36.30 | 2.47 | 17.38 | 7.04 |

low number of *Trichinella* muscle larvae, but their number on day 30 p.i. increased to a comparable value (intoxication with 0.2 mg.kg⁻¹ b. w.) or half-value (intoxication with 1.0 mg.kg⁻¹ b. w.).

A common biological characteristic used to evaluate *Trichinella* species infectivity is the Reproductive Capacity Index (RCI) — the ratio of the total number of larvae recovered from an animal to the number of larvae administered to it. The first is the product of 4 components: the number of females established; their fecundity; their reproductive life span; and the survival of muscle larvae. At the muscle phase of *T. Spiralis* infection, RCI was decreased and markedly reduced by the high intoxication with Cd and Hg (Table 2).

The observed differences in parasite numbers could be explained by the different heavy metal toxicity. Exposure to Pb ions greatly increases rodent susceptibility to parasitic infections [5]. Pb can weaken the resistance and increase the mortality risk in experimental animals and reduce the antibody production [1], [6]. Cadmium has pro-inflammatory properties [14]; already developed inflammation in mice intoxicated with cadmium significantly reduced the number of adult worms. A higher cadmium dose had probably a disturbing effect on their fertility. The effect of heavy metals on mammal reproduction has been confirmed by several studies. Pace et al. [18] reported that exposure to Pb have resulted in a reduced number of testicular macrophages, relating to a significant decline in fertility. The authors suggested that testicular macrophages could be one of the main targets of lead and exposure to even low doses in neonatal and adolescent stages can affect reproductive functions in adult male mice.

Heavy metal toxic effects on helminths has not yet been studied in detail; a single study [2] on the intestinal fish parasites reported the increased occurrence of morphological malformations in the reproductive organs in *Proteocephalus percae* tapeworms, parasitising in perch in heavy metal contaminated environment (the Ružín water reservoir). Though we did not focus on determining the disturbing effect of heavy metals on *Trichinella* reproduction, a higher dose of cadmium significantly reduced the number of muscle larvae when compared to the other experimental groups. Cadmium can disturb the hormonal balance in mammals, e.g. altering the concentration of pituitary gland hormones, such as prolactin, adrenocorticotropine hormone, growth hormone, luteinizing hormone, and follicle stimulating hormone [13]. In helminths, cadmium can also affect their metabolism and internal organ structure.

Exposure to Hg enhanced the sensitivity of organisms to infection due to the inhibition of cell and humoral responses [15]. On the other hand, *in vitro* studies with inorganic mercury referred to a stimulation of lymphatic system activity and humoral immunity and an increase in a number of T and B spleen cells in mercury susceptible mice strains [8], [16]. An acute or chronic mercury intoxication, even its sub-toxic doses, disabled the immune system that is manifested by increased susceptibility to autoimmune disorders or immunosuppression [7], [11], [22].

Environmental contamination can in some cases increase the effects of parasitism, e.g. increase population

density of suitable intermediate hosts and definitive hosts, or negatively affect the host defence mechanisms, thus increasing the host susceptibility (intoxication with Pb) [12]. However, the pollutants can decrease parasitism in such a way that infected hosts suffer more from environmental load (intoxication with Hg) than do infection-free hosts, parasites are more susceptible to pollution than their hosts, or pollution lowers the abundance of essential hosts and intermediate hosts below a threshold for sustained transmission of the parasites. The effect of contaminants may vary among respective parasitic species, and developmental stages [21]. In our experiments, Hg activated adult worms in the small intestine, but reduced the number of muscle larvae.

Many studies have focused on parasite-induced stress in relation to environmental contamination on a physiological level. Such stress is often the first step to weaken the immune response. For instance, fish suffering from parasitic infection are weaker and more susceptible to environmental stress [22]. The unfavourable effects of parasites on the host's health is often camouflaged by an acute toxic effect of the pollutants.

In mice intoxicated with heavy metals and subsequently infected with *T. spiralis*, we observed differences in an intensity of parasitic infection depending on the dose and type of heavy metal. Exposure to cadmium inhibited the development of parasitic infection as early as during the muscle phase. The lead affected the parasite burden in hosts insignificantly; it rather increased the intensity of infection during the muscle phase. Mercury intoxication prolonged the intestinal phase and the larvae remained in the small intestines till day 20 p.i., but that might have been influenced by the poor condition of the mice. Our data refer to the different effect of tested heavy metals on the hosts and intensity of parasitic infection, as well as increased risk of trichinellosis via small rodents in the ecosystems contaminated with heavy metals.

ACKNOWLEDGEMENTS

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CONTENT OF FREE AMINO ACIDS AND SOME PARAMETERS OF THE FUNCTIONAL STATE OF THE LIVER IN BLOOD PLASMA OF HEALTHY AND KETOTIC DAIRY COWS

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ABSTRACT

Metabolic disorders are one of the most important problems in modern cattle production. One of the diseases, which appears to be a major obstacle for an increase in milk productivity is ketosis of dairy cows. Investigations into the changes of the amino acids occurring in the plasma of ketotic cows, allows one to get an insight into the dynamics of this metabolic imbalance, which has significant diagnostic and prognostic value. Therefore, characterizing such changes in the plasma content of free amino acids in ketotic cows are reported in this paper. The study was conducted on 17 Holstein cows in their second to fourth lactations with milk yields above 8000 kg in previous lactations. Nine of these cows were clinically healthy, with negative tests for ketone bodies in their urine, and eight of the cows had positive tests, which also revealed hepatic impairment. Blood samples were taken in March, 2013, two or three weeks after calving. The blood was withdrawn from the jugular vein before the morning feeding. This research has shown that in ketotic cows, the plasma level of ketogenic, aromatic and sulphur-containing amino acids were significantly increased. In addition, the content of glycogenic amino acids and branched-chain amino acids were decreasing, leading to an increase in the ratio of glycogenic/ketogenic amino acids from 4.16 to 6.02. The content of amino acids used in the antioxidant system of the organism decreased on average by 40%, except for the methionine and taurine levels which significantly increased. The ratio between the essential and nonessential amino acids in the plasma of cows affected with ketosis decreased by more than 8%.

Keywords: amino acids; calving; cows; hepatic pathology; ketosis

INTRODUCTION

Ketosis in cows is a severe pathologic process with complex symptoms and signs in which disorders of carbohydrate, lipids and protein oxidation occur. Impairment of functions of the liver, pancreas, other organs and tissues are observed in sick animals [10], [16], [19], [20]. Most commonly the disease is reported in high-yield dairy cows during the first two months of lactation or at the peak of milk production. Amino acids play very important roles in the pathogenesis of ketosis, since they participate in the synthesis of the majority of endogenous biologically active substances, structural proteins, enzymes, some of the hormones, nitrogenous bases, and neurotransmitters [13], [14]. Exogenous amino acids entering the organisms and undergoing oxidation are involved in the energy supply and are subjected to numerous transformations. The metabolism of these substances are controlled by different biochemical and physiological mechanisms, which maintain relatively constant concentrations of amino acids in the blood and tissues. Therefore, the concentration of free amino acids and their derivatives in physiological liquids and tissues may serve as an index of homeostasis, and the pattern of the formation of amino acid reserves in organisms objectively reflect the state of the metabolic balance. The concentration of free amino acids and their derivatives, as such, serve as a control factor for many key metabolic steps. Investigations of

the changes in the plasma content of free amino acids provides the possibility to acquire an insight into the essence of metabolic imbalance in a variety of diseases. Such an approach may contribute to revealing factors of significant diagnostic and prognostic relevance. Reasoning from this fact, the objective of our study was to investigate and to compare the plasma content of free amino acids in healthy and ketotic dairy cows.

MATERIALS AND METHODS

This study was performed in March, 2013, on seventeen Holstein cows in their second to fourth lactations with a milk yield above 8000 kg in previous lactations. Animals were kept on a farm in similar conditions and received the same diet.

Clinical examinations of the cows were performed and the content of ketone bodies in their urine was detected with diagnostic strips (Ketophan, Pliva). During the examinations it was revealed that some animals (n=8) were lying up, revealed rapid weight loss and milk yield, they looked depressed, and some of them had muscular tremor. After contact of the indicator strips with the urine, the colour turned violet, indicating the presence of ketone bodies. During the laboratory investigations, the diagnosis was confirmed by using fluorometric determinations of beta-hydroxybutyrate (BHBA) levels in the blood and milk samples. Animals with positive results (the presence of ketone bodies in the urine) were relegated into a separate group. In order to obtain plasma, blood samples were withdrawn two or three weeks after calving, from the jugular vein before the morning feeding, into sterile tubes with heparin and immediately centrifuged at 3000 rpm. The plasma was frozen at -20°C until analyses were done (up to three month following).

In order to obtain serum, venous blood was withdrawn into sterile plastic tubes, which were left at room temperature for clot formation. After that, the blood clots were separated from the tube walls with a thin glass rod, settled blood serum was transferred into new sterile tubes and centrifuged for 15 minutes at 1000 rpm. This blood serum was again transferred into new sterile tubes and stored in the refrigerator at a temperature from 4 to 6°C up to two days until blood tests were performed. The content of free amino acids was determined in the plasma using the ion-exchange techniques on the amino acid analyser Biotronik LC 6001 (Germany). The serum was used for the determination of: AST and GGT activity (Reitman & Frankel's method and kinetic colorimetric method, respectively [18]); levels of albumin (bromocresol green colorimetric method); and bilirubin (colorimetric method by Jendrasik and Grof [15]), using a biochemical analyser Humalyzer 2000 (Germany) and test-systems manufactured by Human (Germany).

The results were processed statistically in Excel, arithmetical mean value (x) and the standard error of the mean (SEM) was calculated as well as the significance of the difference between two variation series.

RESULTS AND DISCUSSION

The investigation of the amino acid content in the plasma of healthy and sick cows revealed a range of distinctions (Table 1). In particular, we observed a significantly lower alanine

(by 24.9%; $P < 0.001$), arginine (by 20.1%; $P < 0.001$) and asparagines content (by 28.6%; $P < 0.05$) in the plasma of sick cows in comparison with the healthy ones, which provided evidence of an increase in the utilization of these amino acids. For example, in the case when at the beginning of lactation, there was a low glucose content, amino acids which enter the citric acid cycle or convert into pyruvate, may directly transform into glucose. Consequently, amino acid carbon residues comprise from 15 to 35% in the process of gluconeogenesis [6]. Besides, the decrease in arginine level may be explained by its usage in antioxidant protection of the organism. Some available literature data [2], [11], [17] suggest that some amino acids are able to decrease the damaging and pathological effects caused by various oxidative influences. For instance, there is the proven ability of arginine and citrulline to deactivate superoxide anion radicals, which leads to normalization of myocardial activity under the influence of oxidative agents. The data obtained show a lower level of not only arginine, but also citrulline by 45.5% ($P < 0.01$).

Table 1. Plasma content of free amino acids in healthy and ketotic cows, Means \pm SEM

| Amino acid [$\mu\text{mol.l}^{-1}$] | Healthy n = 9 | Ketotic n = 8 | P < |
|--|-------------------|-------------------|-------|
| Alanine | 157.7 \pm 9.63 | 118.4 \pm 4.93 | 0.001 |
| Arginine | 58.3 \pm 2.16 | 46.6 \pm 2.58 | 0.001 |
| Asparagine | 25.2 \pm 1.93 | 18.0 \pm 0.98 | 0.05 |
| Aspartate | 9.4 \pm 1.21 | 11.1 \pm 1.18 | 0.5 |
| Valine | 120.1 \pm 4.58 | 101.5 \pm 5.42 | 0.05 |
| Histidine | 36.4 \pm 3.96 | 21.0 \pm 1.31 | 0.001 |
| Glycine | 402.6 \pm 24.35 | 287.7 \pm 8.21 | 0.01 |
| Glutamate | 62.8 \pm 3.57 | 61.5 \pm 5.20 | 0.5 |
| Glutamine | 102.3 \pm 5.71 | 204.6 \pm 10.77 | 0.001 |
| Isoleucine | 70.5 \pm 6.22 | 66.7 \pm 9.68 | 0.5 |
| Carnosine | 14.7 \pm 2.93 | 10.0 \pm 1.97 | 0.5 |
| Leucine | 48.6 \pm 2.30 | 79.2 \pm 12.98 | 0.001 |
| Lysine | 57.3 \pm 7.63 | 59.8 \pm 6.04 | 0.5 |
| Methionine | 16.8 \pm 1.48 | 24.2 \pm 1.42 | 0.01 |
| Proline | 135.5 \pm 5.21 | 114.5 \pm 22.76 | 0.1 |
| Serine | 56.9 \pm 5.74 | 65.7 \pm 8.93 | 0.5 |
| Taurine | 17.7 \pm 1.35 | 27.02 \pm 2.49 | 0.05 |
| Tyrosine | 27.6 \pm 2.05 | 31.1 \pm 5.64 | 0.5 |
| Threonine | 45.4 \pm 3.85 | 37.9 \pm 9.90 | 0.5 |
| Tryptophan | 38.6 \pm 2.95 | 56.8 \pm 4.40 | 0.01 |
| Phenylalanine | 24.3 \pm 0.53 | 29.8 \pm 2.64 | 0.05 |
| Phosphoserine | 15.9 \pm 5.42 | 24.2 \pm 2.33 | 0.5 |
| Cysteine | 6.64 \pm 0.37 | 1.67 \pm 0.73 | 0.001 |
| Citrulline | 77.4 \pm 6.30 | 42.2 \pm 5.78 | 0.01 |
| 1-methylhistidine | 9.2 \pm 3.97 | 10.9 \pm 0.93 | 0.5 |
| 3-methylhistidine | 6.0 \pm 1.03 | 28.4 \pm 2.06 | 0.001 |

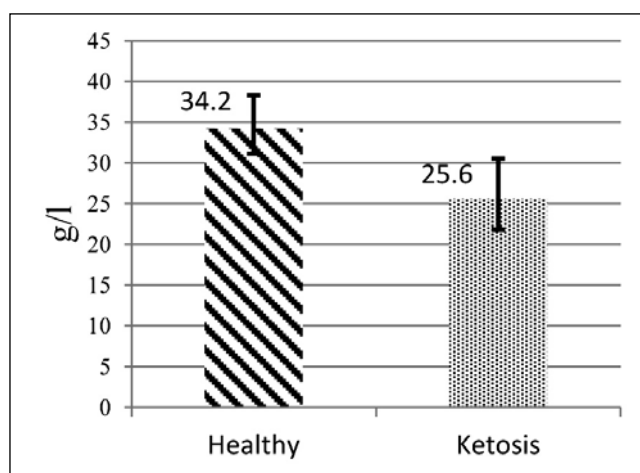


Fig. 1. Serum albumin concentration in cows

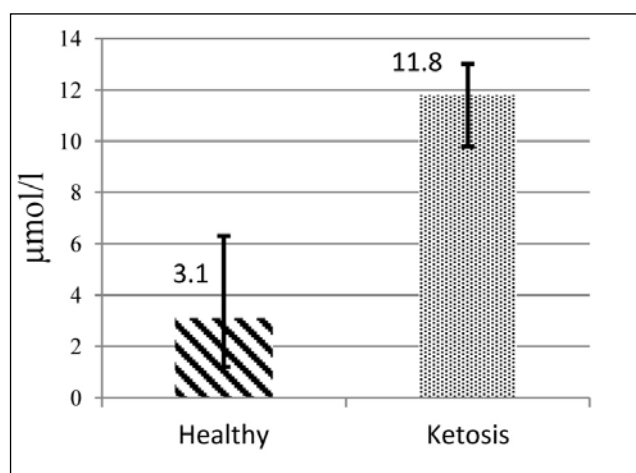


Fig. 2. Serum levels of total bilirubin in cows

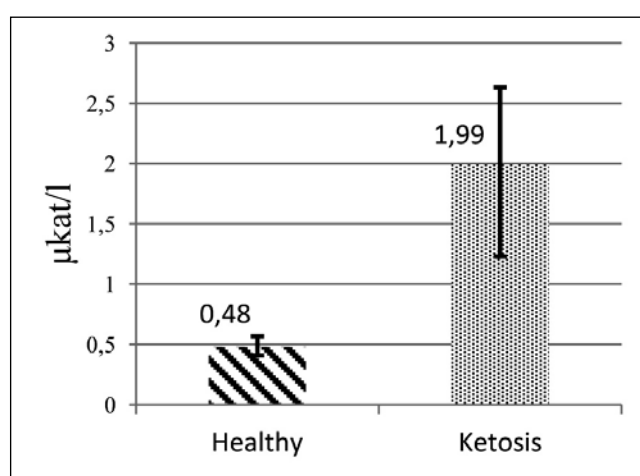


Fig. 3. Serum activity of aspartate aminotransferase in cows

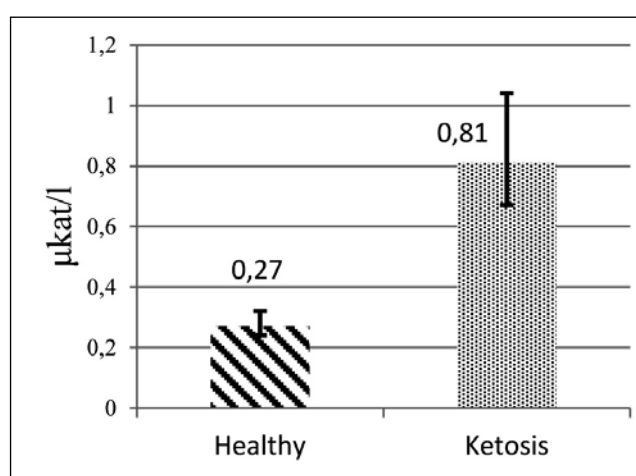


Fig. 4. Serum activity of gamma-glutamyltransferase in cows

In contrast with the glycogenic amino acids, the plasma content of ketogenic amino acids in ketotic cows was higher. Particularly, the content of phenylalanine was higher by 18.5 % ($P < 0.05$), tyrosine by 11.3 %, leucine by 38.6 % ($P < 0.001$) and tryptophan by 32 % ($P < 0.01$). On the one hand, the higher level of amino acids may be related to activation of gluconeogenesis, but then, such changes may be associated with the peculiarities of the degradation of different amino acids.

Aromatic and sulphur-containing amino acids decompose only in the liver. In particular, phenylalanine, tryptophan, tyrosine and methionine belong to this group. Therefore, the higher level of these amino acids maybe explained by liver injury, due to the lowering of the concentration of intracellular enzymes (which inactivate the above mentioned amino acids) [14].

We have established that in sick cows the serum level of albumin was lower (by 25.1 %; $P < 0.01$; Fig. 1) and the level of total bilirubin was higher almost four-fold ($P < 0.001$;

Fig. 2) in comparison with healthy cows. The activities of AST (more than four-fold; $P < 0.001$; Fig. 3) and GGT (three-fold; $P < 0.001$; Fig. 4) were also higher. According to Table 1, in sick animals, the plasma levels of not only phenylalanine and tyrosine were higher, but also of methionine (by 30.6 %; $P < 0.01$).

Disorders of the degradation of aromatic and sulphur-containing amino acids are the cause of excessive formation of endotoxins, such as phenol, mercaptans, and indole, which exacerbate the pathological process. Nevertheless, an increased content of methionine also has a positive aspect, because this amino acid plays a significant role in the system of antioxidant protection. The antioxidant effect of methionine is explained by the fact that it is a good source of sulphur, which inactivates free radicals and enhances the utilization of lipids, preventing their deposition in the liver and vascular walls. The synthesis of taurine and cysteine, which is a precursor of glutathione, depends on the level of methionine [5], [8], [9]. The results show significantly ($P < 0.05$) higher taurine levels (by 34.5 %) in the plasma of ketotic cows, which is probably associated with an increase in the activity of com-

pensatory mechanisms in animal organisms. There are also available data [8] concerning the positive role of proline and histidine in the antioxidant defence of organisms. Proline is an effective quencher of singlet oxygen; it enhances prevention of cell damage and death under oxidative stress. Histidine has the ability to trap peroxy radicals, prevents protein carboxylation and formation of protein cross-links. These effects are probably determined by physico-chemical properties, related to their ability to react with active forms of oxygen. As it can be seen from the results, in sick animals, the content of proline and histidine was lower by 15.5% and by 42.3% ($P < 0.001$) respectively in comparison with healthy cows. Besides, the level of cysteine was lower by 74.8% ($P < 0.001$) and glycine by 28.5% ($P < 0.01$), which might be explained by the usage of these amino acids in the synthesis of glutathione, which, in turn, is utilized for organism detoxification.

This research has shown that the level of branched-chain amino acids (valine and isoleucine) was lowered (by 15.5%; $P < 0.05$ and 5.4%, respectively), except for leucine, which also belongs to branched-chain amino acids, which exerts a ketogenic action. These three essential amino acids are degraded mostly in muscle tissue and play a significant role in energy metabolism, particularly in the synthesis and deposition of glycogen. Consequently, an imbalance in amino acid content in the blood occurs in ketotic cows, which is an unfavourable factor in the course of the disease.

Besides the changes in plasma amino acid content, we should also draw attention to the significantly higher levels of glutamine (by 50%; $P < 0.001$) and 3-methylhistidine (by 78.9%; $P < 0.001$) levels. These increased glutamine levels may contribute to the growth of glucocorticoid activity, which leads to an increase of glutamine synthetase activity, the enzyme which catalyzes the formation of glutamine from glutamic acid and ammonia. High level of 3-methylhistidine may be considered as a marker of catabolism intensification in muscle tissue. It is an amino acid specific for contractile proteins (actin and myosin). During degradation of these proteins the given amino acid enters the blood circulation, but due to the absence of specific tRNA, it is not used for protein synthesis, nor metabolized and in an unchanged form is excreted in the urine [1], [3], [4], [7], [12], [21]. These properties make the determination of 3-methylhistidine content, an informative indicator of muscle protein catabolism.

The investigation of the ratio between glycogenic amino acids (alanine, arginine, asparagine, valine, histidine, glycine, glutamine, methionine, proline, serine, threonine, cysteine) and amino acids with ketogenic (leucine, lysine) or mixed action (tyrosine, tryptophan, phenylalanine) becomes also noteworthy. Thus, in the plasma of ketotic cows, this ratio is higher by 30.9% (4.16 vs. 6.02), and the ratio between essential and nonessential amino acids is lower by more than 8% (from 1.81 vs. 1.66).

CONCLUSIONS

Ketotic cows exhibit an imbalance of amino acid content in their blood. This is an unfavourable factor in the course

of the disease. Particularly, in comparison with healthy animals, in ketotic cows, the content of glycogenic acids is significantly lower, and the level of ketogenic amino acids is higher, which in turn causes an increase in the ratio of glycogenic/ketogenic amino acids from 4.16 to 6.02.

The content of amino acids, which are used in the antioxidant defence of organisms (arginine, citrulline, proline, histidine, cysteine and glycine), were lower, on average by 40%, except for methionine and taurine, the content of which was significantly higher.

Impairment of hepatic function in ketotic cows causes elevation of the levels of aromatic and sulphur-containing amino acids and lowering of the content of branched-chain amino acids. The ratio of essential and nonessential amino acids in the plasma of cows affected with ketosis was lower by more than 8% (1.81 vs. 1.66).

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ASSESSMENT OF SELECTED ANTIOXIDANTS AFTER FERROCENYL-CHALCONES TREATMENT OF MITOCHONDRIA

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ABSTRACT

Ferrocene related organometallic compounds are promising compounds for the design of novel drugs as they may exhibit much greater activity in living systems than the original compound without the ferrocene moiety. We investigated the activities of enzymatic and non-enzymatic antioxidant levels of five ferrocenyl derivatives at a final concentration of 0.1 and 0.01 $\mu\text{g} \cdot \text{ml}^{-1}$ in mitochondria (the smallest, sufficiently closed system for the study of oxidative damage *in vitro*). The highest ability to influence the redox state was demonstrated by the ferrocenyl chalcone, 1,1'-bis [(E)-(2-indanonyl)-methylidene] ferrocene, which decreased glutathione peroxidase activities and caused glutathione depletion, when used in a final concentration of 0.01 $\mu\text{g} \cdot \text{ml}^{-1}$. Similarly, a greater ability to cause oxidative damage contributing to retrograde redox signalling from the mitochondria to the cytosol and nucleus, could result from the treatment with 1,1'-bis [(E)-(2-tetralonyl)-methylidene] ferrocene. However, further investigations of their biological activity are necessary.

Key words: antioxidant; ferrocene; chalcone; metallocene

INTRODUCTION

Ferrocene (Fc), as an effective bioisostere, is becoming an applicable platform for drug design by virtue of its redox properties, high lipophilicity and three dimensional metallocene unit. There-

fore, ferrocene moieties may lead to some positive changes in selectivity toward biological targets compared to phenyl or alkyl groups (15). As aromatic compounds, they show interesting redox properties and chemical stability, and several of their derivatives may also affect their relative safety in many mammalian species. Many ferrocenyl compounds display interesting cytotoxic, antitumor, antimalarial, antifungal and DNA cleaving activities [17]. Ferrocene can act; as one electron donor undergoing oxidation to the ferricenium ion (Fe^{3+}) or, as a source of reactive oxygen species (ROS) that has a direct influence on its activity [12], [16]. The combination of a ferrocenyl moiety with heterocyclic structures may increase biological activities or create new medicinal properties.

Chalcones are naturally occurring derivatives of the parent compound 1,3-diphenyl-2-propen-1-one and belong to the flavonoid family of organic compounds. Chalcones have been reported to have a broad range of biological activities, such as: antimalarial, antibacterial, antitumour, antihyperglycemic and anti-HIV. A ferrocenyl chalcone (Fc-chalcone) derivative is formed when either ring A or B is replaced with a ferrocenyl group [1]. Series A and B compounds differ significantly in their physicochemical properties. The series A compounds have been characterized by; a higher resistance to oxidation and generally lower lipophilicity, but more polarized carbonyl bonds, than in series B [17]. When ferrocene is ring B, free radical generation should be greater, since ferrocene is more readily oxidized to ferricenium. Since the oxidizability of ferrocene can largely influence the oxidizing properties of Fc-chalcones, there is evidence supporting the information to what extent they are able to scavenge radicals, and to what extent they are able

to donate electrons to other suitable acceptors with the formation of free radicals and subsequent biological response.

We investigated the effects of five ferrocenyl derivatives (of which four were ferrocenyl chalcones) at two concentrations which most influenced the viability of cancer cell lines, on the activity of enzymatic antioxidants and levels of non-enzymatic antioxidants in mitochondria (which are the smallest and sufficiently closed system for the study of the potential effects on cellular metabolism *in vitro*).

MATERIALS AND METHODS

Compounds; A: (E)-3-(ferrocenemethylene)-4-chromanone; B: (E)-2-(ferrocenemethylene)-cyclohexanone; C: 1,1'-bis [(E)-(2-indanonyl)-methylidene] ferrocene; D: 1,1'-bis [(E)-(2-tetralonyl)-methylidene] ferrocene; and E: 2-cyano-3-ferrocenylacrylonitrile; were synthesized and purified as described previously [4, 13]. Their structures were characterized by IR and NMR spectroscopy. The purity of the compounds was checked using thin layer chromatography and gas chromatography. All other chemicals used were of analytical grade and, if not stated otherwise, purchased from Sigma — Aldrich or Serva (Heidelberg, Germany). Liver mitochondria were isolated using the method of Fernández-Vizar et al. [5]. Measurements were carried out in a respiratory medium (80 mmol.l⁻¹ KCl, 300 mmol.l⁻¹ KH₂PO₄, 300 mmol.l⁻¹ K₂HPO₄, 15 mmol.l⁻¹ TRIS-HCl, 6 mmol.l⁻¹ MgCl₂, and 0.78 mmol.l⁻¹ EDTA) at pH 7.4. The tested compounds were added in a final concentration of 0.1 µg.ml⁻¹ and 0.01 µg.ml⁻¹. The activity of glutathione reductase (GR; EC 1.6.4.2) was measured according to a modified method described by Carlberg and Mannervik [2]; that of glutathione peroxidase (GPx, EC 1.11.1.9) was measured as described by Flohe and Gunzler [6] and that of superoxide dismutase (SOD, EC 1.15.1.1) by means of the SOD-Assay Kit-WST (Fluka, Japan), following the user manual provided. The reduced glutathione (GSH) content was measured by the modified method of Floreani et al. [7] using Ellman's reagent. All the measured parameters were calculated per mg or g of mitochondrial protein, determined by using the bicinchoninic acid assay.

The results were expressed as the mean ±SD of three independent measurements. The significance of differences between groups was determined using an unpaired Student *t*-test and were considered significant at: * — *P* < 0.05; ** — *P* < 0.01; *** — *P* < 0.001.

RESULTS AND DISCUSSION

The production of ROS by mammalian mitochondria is important because it underlies pathology caused by oxidative damage and contributes to retrograde redox signalling from the organelle to the cytosol and nucleus [10]. Mitochondria generate approximately 2–3 nmol of O₂⁻ per mg of protein, the ubiquitous presence of which, indicates it is to be the most important physiological source of this radical in living organisms [9]. Therefore, it is important to understand O₂⁻ production within isolated mitochondria under conditions that mimic those that may arise *in vivo* under physiological or pathological conditions. A constraint of us-

ing isolated mitochondria is that the system is complicated and difficult to manipulate [3]. O₂⁻ undergoes a dismutation reaction to H₂O₂ spontaneously, or is accelerated by about four orders by SOD, in the mitochondrial matrix by Mn-SOD, and in the intermembrane space by Cu, Zn-SOD [8]. The activity of SOD was decreased (compared to the control) in the mitochondria with higher concentrations of the tested compounds, significantly by the treatment with the first three ferrocenyl chalcone compounds, and also in group E (Fig. 1). Treatment with a lower concentration of the tested substances led contrariwise to a significant increase in the activity of SOD compared to the control, except for compound B which demonstrated a significantly lower activity than the control.

SOD contributes significantly to protecting the organism from the toxic effects of O₂⁻ but its activities produce H₂O₂ from O₂⁻, both at the inter-membrane space and in the mitochondrial matrix. H₂O₂ diffuses rapidly through membranes, and the release of H₂O₂ from the mitochondria to the cytosol reflects the balance between H₂O₂ production and consumption reactions; with the latter mainly involving the reduction of H₂O₂ to H₂O via GPx. The activities of GPx significantly decreased compared with the control under the treatment with both concentrations (Fig. 2), with the sole exception of the treatment with compound E at the final concentration of 0.1 µg.ml⁻¹. However, the differences were not significant at the highest level (*** — *P* < 0.001) when compared to the control. Considering the inability of O₂⁻ to cross the membranes and the decreased requirements on its dismutation according to the measured SOD activities (Fig. 1, using concentration 0.1 µg.ml⁻¹), a decrease in GPx activity (Fig. 2, using concentration 0.1 µg.ml⁻¹) suggested lower peroxide formation and thus lower need for its elimination. However, at relatively high concentrations of H₂O₂ and other hydroperoxides, peroxidases undergo substrate inactivation by the modification of the prosthetic haem group [11]. This is still necessary to consider if we want to explain the high activity of SOD and also significantly reduced activity of GPx (Fig. 1 and 2, using the concentration of 0.01 µg.ml⁻¹).

The activity of GPx is affected by the presence of another antioxidant enzyme, GR, which continuously recycles the oxidized glutathione to the reduced state. In our experiments, GR activities were not significantly changed compared to the controls (Fig. 3). Exposed thiols are present in mitochondria at a high concentration [14]. The regeneration of GSH from glutathione disulphide (GSSG) trapped inside the mitochondria requires GR, which harnesses the more negative reduction potential of NADPH in the process. GR continually recycles the oxidized thiol back to its reduced form, since the potential for GSH regeneration within the mitochondrial matrix is proportionately responsible for the constant degradation of ROS. Another point to be considered is that mitochondrial GSH is not synthesized in the matrix but is imported from the cytoplasmic compartment [14]. Thus, a variation in the mitochondrial matrix GSH ratio could not be balanced by the import. However, in the case of mitochondria, GSH reflected a significant decline in the response to a change in the GSH/GSSG ratio only un-

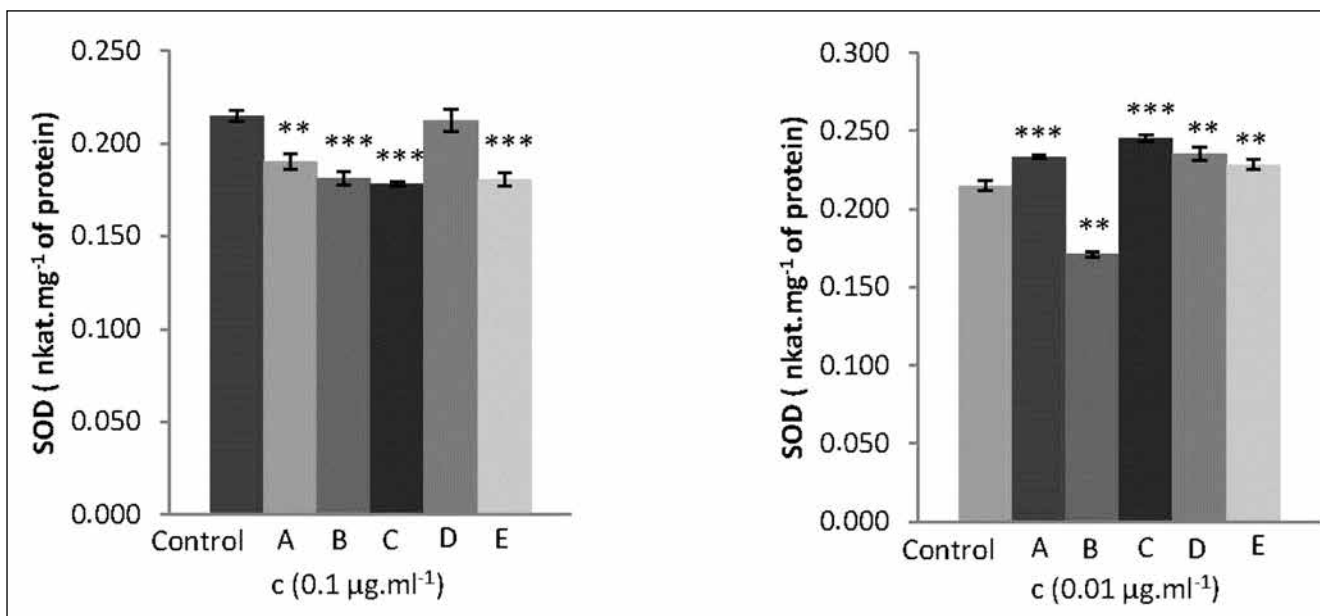


Fig. 1. Superoxide dismutase activities in mitochondria after treatment with the tested compounds
A: (E)-3-(ferrocenemethylene)-4-chromanone; **B:** (E)-2-(ferrocenemethylene)-cyclohexanone; **C:** 1,1'-bis [(E)-(2-indanonyl)-methylidene] ferrocene; **D:** 1,1'-bis [(E)-(2-tetralonyl)-methylidene] ferrocene; **E:** 2-cyano-3-ferrocenylacrylonitrile at the final concentrations 0.1 µg.ml⁻¹ and 0.01 µg.ml⁻¹
 * — Indicates significance of differences; ** — P < 0.01; *** — P < 0.001

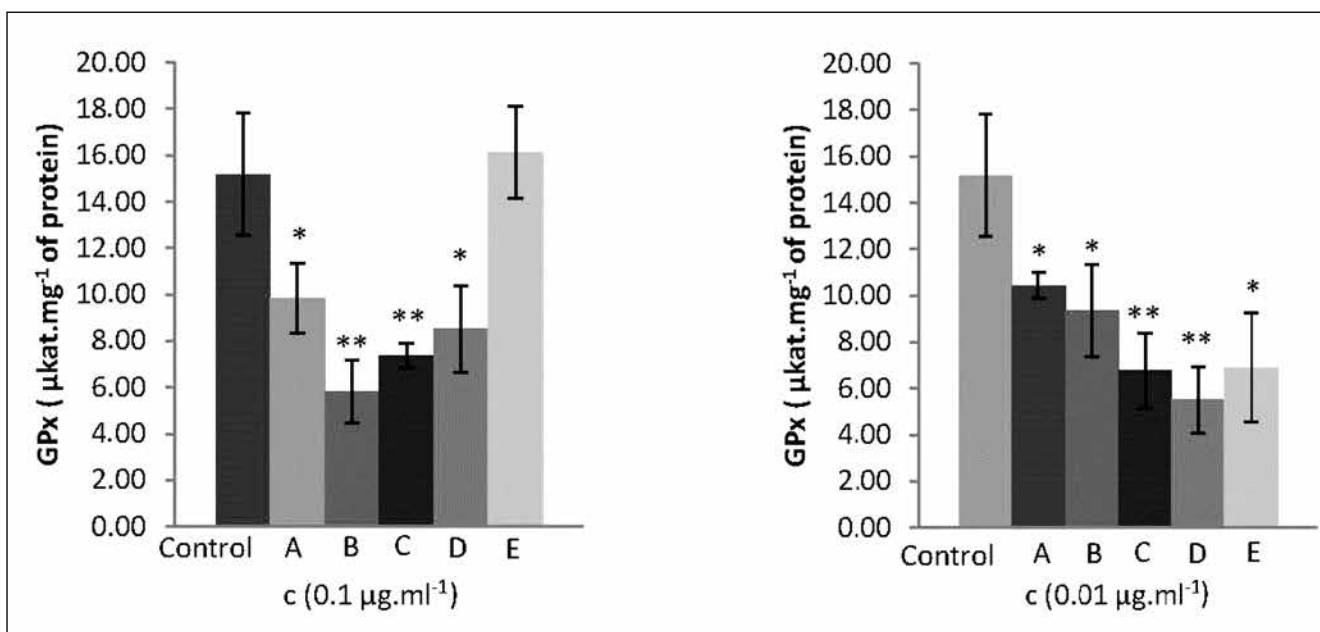


Fig. 2. Glutathione peroxidase activities in mitochondria after treatment with the tested compounds
A: (E)-3-(ferrocenemethylene)-4-chromanone; **B:** (E)-2-(ferrocenemethylene)-cyclohexanone; **C:** 1,1'-bis [(E)-(2-indanonyl)-methylidene] ferrocene; **D:** 1,1'-bis [(E)-(2-tetralonyl)-methylidene] ferrocene; **E:** 2-cyano-3-ferrocenylacrylonitrile with final concentrations 0.1 µg.ml⁻¹ and 0.01 µg.ml⁻¹
 * — Indicates significance of differences; * — P < 0.05; ** — P < 0.01)

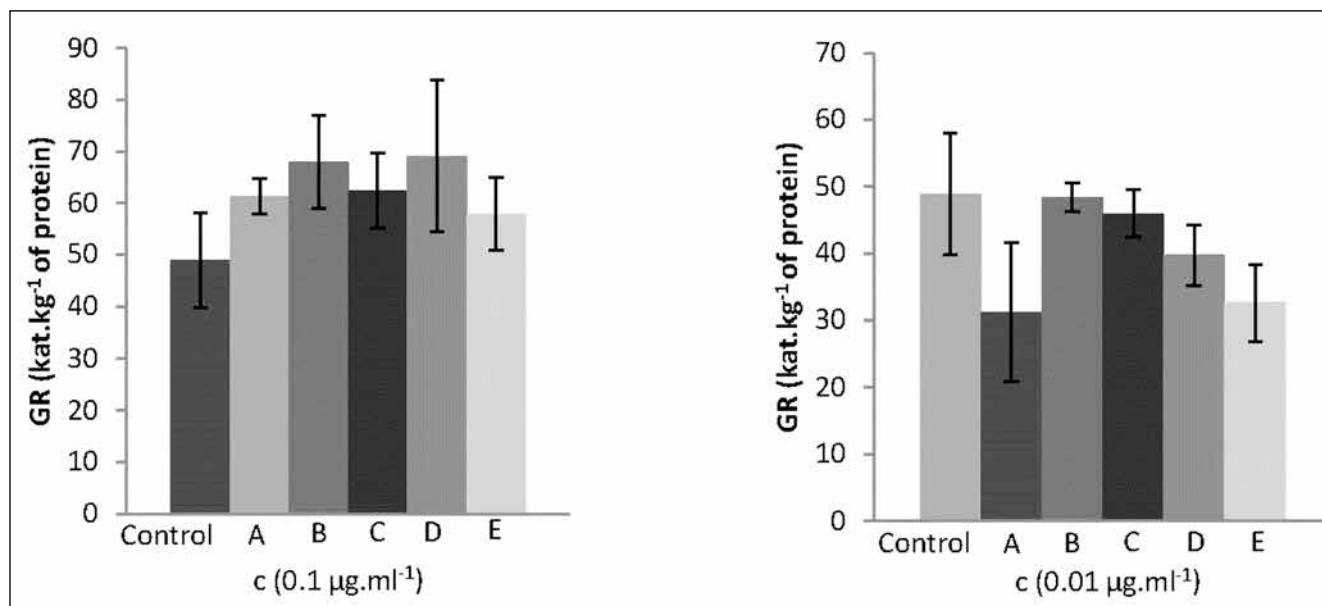


Fig. 3. Glutathione reductase activities in mitochondria after treatment with the tested compounds
 A: (E)-3-(ferrocenemethylene)-4-chromanone; B: (E)-2-(ferrocenemethylene) -cyclohexanone; C: 1,1'-bis [(E)-(2-indanonyl)-methylidene] ferrocene; D: 1,1'-bis [(E)-(2-tetralonyl)-methylidene] ferrocene; E: 2-cyano-3-ferrocenylacrylonitrile at final concentrations 0.1 µg.ml⁻¹ and 0.01 µg.ml⁻¹

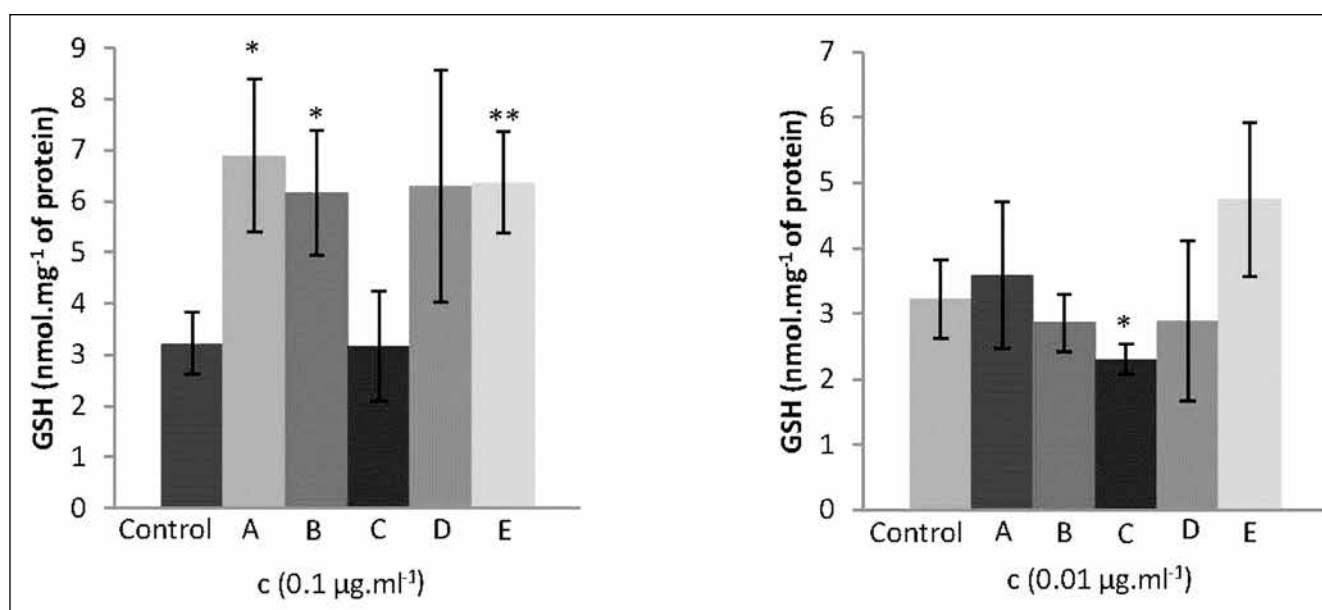


Fig. 4. Levels of reduced glutathione in mitochondria after treatment with the tested compounds
 A: (E)-3-(ferrocenemethylene)-4-chromanone; B: (E)-2-(ferrocenemethylene) -cyclohexanone; C: 1,1'-bis [(E)-(2-indanonyl)-methylidene] ferrocene; D: 1,1'-bis [(E)-(2-tetralonyl)-methylidene] ferrocene; E: 2-cyano-3-ferrocenylacrylonitrile, at final concentrations 0.1 µg.ml⁻¹ and 0.01 µg.ml⁻¹. *Indicates significance of difference
 * — P < 0.05; ** — P < 0.01)

der treatment with compound C in the final concentration of $0.01 \mu\text{g} \cdot \text{ml}^{-1}$ (Fig. 4). Treatment with the higher concentration ($0.1 \mu\text{g} \cdot \text{ml}^{-1}$) led in most cases (A, B, E) to increased levels of reduced glutathione compared to the control.

Ferrocene related organometallic compounds are known and are promising compounds for the design of novel drugs. By attaching organic groups, which creates a conjugated or non-conjugated system, they may exhibit much greater activity in living systems than the original compounds without the ferrocene moiety. Therefore, it would be very interesting to observe their effect on redox homeostasis through the activities of primary antioxidant enzymes and levels of reduced glutathione in the mitochondria. The results indicate that at a concentration of $0.01 \mu\text{g} \cdot \text{ml}^{-1}$, there was a significant increase in SOD activity compared to the control (except with compound B), which itself indicates a high production of superoxide radicals, and thus oxidative stress conditions. Increased dismutation leads to increased formation of peroxides, but a high level is likely to cause inhibition of the catalytic activity of GPx and therefore no significant needs of GR activity on the feedback reduction of oxidized glutathione. GSH is a strong reductant and is able to react not only with peroxides, which can explain the decrease in the measured levels not only when comparing the two final concentrations but especially with compound C, where its level was significantly reduced.

CONCLUSIONS

We observed the most significant effect of concomitant antioxidant enzyme activities together with GSH levels in the case of C derivative, suggesting that 1,1'-bis [(E)-(2-indanonyl)-methylidene] ferrocene demonstrated the highest ability to influence the redox state of cells. The ability to cause oxidative damage, contributing to retrograde redox signalling from the mitochondria to the cytosol and nucleus, was expected for the treatment with 1,1'-bis [(E)-(2-tetralonyl)-methylidene] ferrocene. However, additional analysis of the pro-oxidant, antioxidant and biological activities of this compound is necessary. The low effective concentration of the compound is very promising.

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THE USE OF THE HPLC METHOD IN CONFIRMATION OF COCCIDIOSTAT RESIDUES IN FOOD MATRICES A REVIEW

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ABSTRACT

High Performance Liquid Chromatography (HPLC) is the analytical separation method most frequently used in the detection of residues of pharmacologically active substances in food matrices. It is also utilized to identify coccidiostats in feed and the residues of food. HPLC can be used for both screening and confirmation. These screening and confirmation methods have been validated due to the progress in the field of analytical chemistry. The requirements of the confirmation analysis and ensuring the quality of results are currently met only by liquid chromatography using the procedures consisting of appropriate combinations of cleaning, chromatographic separation, and spectrometry detection.

Key words: coccidiostats; confirmation; development; HPLC

COCCIDIOSIS AND COCCIDIOSTATS

Food producing animals are often affected by the parasitic disease, coccidiosis [40]. Coccidiosis is a severe infectious disease caused by the microscopic parasite of the *Eimeria* species affecting many types of farm animals, especially poultry, in which it causes severe intestinal damage [32], [37], [46]. Even small damage to the intestinal wall can lead to the deteriorated growth of an animal, reduced feed conversion, poor absorption of nutrients from the feed, dehydration, and loss of blood, thus reducing the viability of an

animal [37], [46], [60]. Due to intensive poultry breeding in small concentrated areas in humid and warm environments, this disease spreads very rapidly [37]. The acute form of coccidiosis can cause high mortality, which is one of the main problems in poultry production [11].

In modern poultry production, it is very important to use coccidiostats for the prevention and therapy of coccidiosis. Coccidiostats can be divided into two main groups, i.e. chemical (synthetic) and ionophore (biosynthetic). Synthetic coccidiostats include a wide range of compounds with various chemical structures which are used as feed additives in animal nutrition (halofuginone, robenidine, nicarbazin, diclazuril, decoquinate, and other), or are intended for the therapy of coccidiosis (sulphonamides). Biosynthetic coccidiostats are the substances containing the polyether group and are naturally produced by the fermentation of microorganisms such as *Streptomyces* species and *Actinomadura* species (monensin, narasin, salinomycin, lasalocid, maduramycin, and semduramycin) (Table 1). According to the current applicable legislation, they are only used as feed additives for animal nutrition for coccidiosis prevention [11].

RISKS RESULTING FROM THE USE OF COCCIDIOSTATS AND THE LEGAL ASPECTS

Because of the frequent and long use of coccidiostats in poultry breeding, there is a high risk of residues in food matrices [37]. The careful administration of coccidiostats to poultry is critical to en-

Table 1. Overview of synthetic and biosynthetic coccidiostats

| Synthetic coccidiostats | Biosynthetic coccidiostats |
|-------------------------|----------------------------|
| Halofuginone | Monensin |
| Robenidine | Narasin |
| Nicarbazin | Salinomycin |
| Diclazuril | Lasalocid |
| Decoquinate | Maduramycin |
| and others | Semduramycin |

sure reduced residues of these substances in animal products made from poultry, in order to minimization general health risks [42]. The residues of coccidiostats can be present in the food of animal origin due to: an improper dose of coccidiostats; improper administration of the medicated feed during certain periods of time; failure to adhere to the withdrawal period; or due to the cross-contamination in feed production while using the same production line (37). If the total exposure in animals does not exceed the allowable daily limit for each coccidiostat, then the contents of each coccidiostat residue in edible animal tissues remain sufficiently low. The most frequently detected coccidiostats in poultry products include: robenidine, nikarbazin, diclazuril, and lasalocid. The major reasons for the presence of their residues are the improper use of coccidiostats and cross-contamination of the feed [23].

Biosynthetic coccidiostats are well-known for their cardiovascular effects. The majority of these effects have been characterized for monensin, as a model substance for the entire group of the above mentioned coccidiostats. Their action is based on increasing the coronary blood flow with the subsequent dilatation of the coronary arteries [27]. Biosynthetic coccidiostats are also the cause of muscular degeneration and have a toxic effect on the nervous system that leads to neuropathy, manifested by myelin degeneration and ataxia [38]. The adverse effects of synthetic coccidiostats on humans as consumers have not been described, except for residues of sulphonamides that can cause severe damage to the consumer's health. Sulphonamides are well-known for their negative effects on the thyroid in connection with the development of thyroidal carcinoma [42].

REGULATIONS APPLYING TO COCCIDIOSTATS

For the purpose of human health protection, the following regulations apply to coccidiostats:

Regulation (EC) No.1831/2003 of the European Parliament and of the Council on Additives for Use in Animal Nutrition. The regulation currently authorises eleven coccidiostats as the supplementary substances intended for broilers, turkeys, and rabbits (monensin, narasin, salinomycin, lasalocid, maduramycin, semduramycin, robenidine, halofuginone, nikarbazin, diclazuril, and decoquinate) [53].

Regulation of the Government of the Slovak Republic No. 320/2003 Coll. on Monitoring of Certain Substances and their Residues in Live Animals and in Products of Animal Origin. The governmental regulation determines the obligatory inspection of coccidiostat residues in beef cattle, sheep, goats, swine, equidae, poultry, in eggs, in rabbit meat, game, in animals from farm breeding and in feed [51].

Commission Regulation (EU) No.37/2010 on pharmacologically active substances and their classification regarding maximum residue limits in foodstuffs of animal origin. The applicable wording of the regulation determines the maximum limits for residues of; amprolium, diclazuril, decoquinate, halofuginone, monensin, lasalocid, sulphonamides, and toltrazuril in milk, meat, liver, kidneys, eggs, skin, and fat. The maximum residue limits have been determined in compliance with the generally acknowledged principles for safety assessment, considering the toxicological risks, environmental pollution, as well as microbiological and pharmacological effects of the residues [17].

Commission Regulation (EC) No. 124/2009, setting maximum levels for the presence of coccidiostats or histomonostats in food resulting from the unavoidable carry-over of these substances in non-target feed. The applicable wording of the regulation determines the maximum contents of; monensin, narasin, salinomycin, lasalocid, maduramycin, semduramycin, robenidine, halofuginone, nikarbazin, diclazuril, and decoquinate in milk, liver, kidneys, eggs, skin, fat, and in other food products [16].

Commission Decision No. 2002/657/EC, implementing Council Directive 96/23/EC concerning the performance of analytical methods and the interpretation of results. The decision stipulates the rules for analytical methods to be used in the testing of the official samples and especially the general rules for interpretation of analytic results of official control of these samples [15].

CHROMATOGRAPHIC SEPARATION TECHNIQUES USED IN CONFIRMATION OF COCCIDIOSTAT RESIDUES

With regard to the presence of coccidiostat residues in the food of animal origin, it is necessary to have certain techniques established in practice for their identification and quantification. Literature sources describe plenty of methods for the analysis of coccidiostat residues in food matrices, such as liver, meat, eggs [25], [41], [43], [50], [60], [61] and milk [48].

Certified laboratories perform the routine inspection of coccidiostats which requires sensitive and reliable methods. The majority of the methods developed for the identification of coccidiostat residues in food matrices are based on the liquid chromatography (LC) with spectrophotometry or fluorescence detection, but also with the mass spectrometry, especially while using the tandem mass spectrometry, enabling sensitive identification and quantification of the target compound [11].

THIN LAYER CHROMATOGRAPHY (TLC)

In the early 1970's, the thin layer chromatography (TLC) was used for the qualitative detection of banned substances (thyreostat-

ics and some anabolics) [20]. This technique was also the first fast semi-quantitative analytical method developed also for identification of coccidiostat residues which was based on the division of individual substances between the ascending mobile phase and the stationary phase of the thin layer. In the TLC method, identification of analytes is carried out either directly under the ultraviolet (UV) radiation emitting lamp (UV lamp), or by spraying a plate with a suitable reagent which forms, together with analytes, products visible with the naked eye.

One of the ways how to increase the sensitivity and specific nature of this method is to combine it with the microbiological detection (TLC/B), i.e. with the bacterial strain [59]. Microbiological detection is generally used to identify the antimicrobial activity of pharmacologically active substances. The detection is carried out directly on the chromatographic plate placed on the surface of agar containing the respective bacterial strain that is incubated under the prescribed conditions. The separation and microbial detection is thus carried out on the same plate. The visualisation of zones is usually carried out using the dehydrogenase-active detection reagents [13].

In 1967, Donoho et al. [22] first described the identification of residues of biosynthetic coccidiostats using the TLC with microbiological detection using the bacterial strain *Bacillus subtilis*. Vanderkop et al. [65] gradually enhanced this method by improving the sample preparation, the sample cleaning process, agar, and solvents. Coccidiostat residues can be identified using also other bacterial strains, such as *Bacillus stearothermophilus*, which was used to identify residues of salinomycin [34], or residues of monensin, salinomycin, and lasalocid [66]. The available literature indicates that the TLC was used in a smaller extent also for the identification of synthetic coccidiostats, for example robenidine in the feed [8].

Asukabe et al. [3] detected monensin, lasalocid, and salinomycin using the high-performance thin layer chromatography (HPTLC). The HPTLC principle is the same as in the TLC. The differences are: for example, in the thickness of the chromatographic layer of the plate; in the number of theoretical compartments on a plate; in the volume of the sample dose; etc. The HPTLC was successfully used for the qualitative and quantitative detection of residues of sulphonamides in food matrices, e.g. meat.

The application of this TLC method in practice, while identifying coccidiostat residues, is less frequently used, due to the extended use of other procedures based on the use of high-performance liquid chromatography as the separation technique [64]. At present, the TLC method is used exclusively for the screening purposes [69].

HIGH-PERFORMANCE LIQUID CHROMATOGRAPHY WITH SPECTROPHOTOMETRIC DETECTION (HPLC-UV) AND FLUORESCENCE DETECTION (HPLC-FLD)

History shows that the largest number of confirmation methods described for the analysis of coccidiostat residues in food matrices and in feed, was represented by the analytical separation method, the high-performance liquid chromatography (HPLC). The specific nature of this method, however, depends on the detector used which also affects the selectivity and sensitivity of the method [63].

The use of the UV detector in the analysis of residues is limited,

as this detector is only suitable for identification of those analytes which show absorbance within the wave length range from 180–600 nm. Similarly, the use of FLD detector is limited by the ability of analytes to excite at the specific wave length and then emit the radiation of a specific wave length [29], [59].

Some analytes cannot be detected by absorbance in the UV area or fluorescence without further adjustment, e.g. derivatization (insertion of a suitable chromophore/fluorophore into their structure). Thus, it requires the reaction with a selected agent, such as 9-antryldiazomethane (ADAM), dansyl chloride, 2,4-dinitrophenylhydrazine, 1-(bromoacetyl) pyrene, vanillin, or dimethylaminobenzaldehyde. If the reaction runs before the injection into the HPLC system, it is called the pre-column derivatization; if it runs after the chromatographic separation on the column, it is called the post-column derivatization [59]. The chromophore is a functional group in the substance molecule that absorbs the UV radiation in the area; for example, a molecule with conjugated double bond, carbonyl, amino and aromatic group.

Identification of some coccidiostat residues, e.g. halofuginone, robenidine, nicarbazin, diclazuril, and lasalocid, using the HPLC-UV technique, can be carried out directly without any chemical analyte pre-adjustment. The identification of biosynthetic coccidiostats cannot be carried out directly, as there is no chromophore in their structure. The HPLC-FLD technique can be used without the chemical pre-adjustment of analyte only in lasalocid identification. Other coccidiostats are only detectable using this technique after they are derivatised [59], [63].

The HPLC with the UV detection has been used since the middle of the 1970's, compared to the HPLC with the FLD detection, the use of which began a little later [20]. The use of the HPLC with the UV detection method is described in the identification of: diclazuril in feed [21]; nikarbazin in tissues, eggs, and in feed [24]; monensin, narasin, and salinomycin in tissues [31]; nikarbazin in feed [26]; monensin in poultry tissues [43]; and robenidine in tissues [73]. Robenidine identification in feed was described also by the method of the high-performance liquid chromatography with the diode array detector (HPLC-DAD) which also provides the information on the spectrum of analyte in the UV area, which is important for coccidiostats identification [39]. The HPLC with the FLD detection was developed for: lasalocid in liver [70]; monensin, narasin, and salinomycin in liver [45]; monensin, salinomycin, and lasalocid in feed [2]; and robenidine in tissues [14].

GAS CHROMATOGRAPHY WITH MASS SPECTROMETRY (GC-MS)

In the 1990's, the gas chromatography with mass spectrometry (GC-MS) appeared on the market for a reasonable price [20]. Although the gas chromatography is able to separate the compounds with a wide range of chemical and physical properties, for the coccidiostat residues, it lacks the high resolution efficiency [54]. One disadvantage of this method is that in the case of identification of coccidiostats, it is time consuming due to difficult sample adjustment before the analyse [57], and considering the fact that it works at high temperatures, this method is less suitable for the analysis of coccidiostat residues due to their low volatility and thermal instability [27], [59].

The GC-MS method was used to identify both, synthetic and biosynthetic coccidiostats, such as halofuginone in the feed [1], lasalocid in the liver [71], monensin in the meat [58], and diclazuril in the feed [5].

LIQUID CHROMATOGRAPHY WITH MASS SPECTROMETRY (LC-MS)

Since the late 1990's, the liquid chromatography with the mass spectrometry (LC-MS) belongs to the standard equipment in laboratories [20]. The efficiency of the mass spectrometry lies in the fact that by means of identification of precursor ions and products of its fragmentation, it provides the information on the analyte structure [33]. The most frequently used mass analyser, is the simple quadrupole that represents a standard analyser for both, gas and liquid chromatography, with sufficient resolution (i. e. the analyser's ability to provide differentiated signals for ions of similar mass and charge — m/z). It became frequently used for analyses thanks to its simplicity and low price.

The review of the available literature indicates that the LC-MS method was used to detect residues of biosynthetic coccidiostats in the food of animal origin [7], [19], [33], nikarbazin in tissues [44], [47], and in eggs [6], robenidine in feed [39], [47], diclazuril in plasma [18] etc.

Schneider et al. [55] was the first team to apply the LC-MS electrospray ionization technique to identify the residues of biosynthetic coccidiostats in the liver of broiler chickens.

Today, the mass spectrometry represents the basic confirmation method in certified laboratories, and in a combination with the liquid chromatography it is a suitable option for identification and quantification of coccidiostat residues in food and in feed [4], [12].

LIQUID CHROMATOGRAPHY WITH TANDEM MASS SPECTROMETRY (LC-MS/MS)

By concurrent engaging of several mass analysers, a highly sensitive and selective tandem mass analyser is formed [59]. For analyte quantification, the so-called triple quadrupole (QQQ) is most frequently used. It consists of two mass analysers of the quadrupole type and the collision cell, and it is also the most frequently used mass spectrometer for identification of coccidiostat residues in food matrices, such as milk, muscles, liver, kidneys, eggs, and in feed. In certified laboratories, however, residue analyses are gradually carried out preferably using other mass analysers, such as the ion trap (IT) that appeared sometimes in 1990 and later the high-resolution mass spectrometers, such as the time-of-flight analyser (TOF) [4]. Innovations in the field of liquid chromatography are represented by the combinations of quadrupole either with the ion trap, or with the time-of-flight analyser. Also well-known, is the combination of quadrupole, where the collision cell and the third mass analyser are replaced with the ion trap and then followed by the time-of-flight analyzer (QITTOF) [30].

For identification of biosynthetic coccidiostats (monensin, narasin, salinomycin, and lasalocid) in the feed for poultry, Wang et al. [68] used the matrix-assisted laser desorption/ionization technique combined with the time-of-flight mass spectrom-

etry (MALDI-TOF-MS). Other well-known studies describing the identification of synthetic and biosynthetic coccidiostats by using the LC-MS/MS technique are: 28, 10, 48, 49, 52, 56, 60, 62, 67 and many others.

Newer more sophisticated techniques are nowadays represented by linear ion traps (LIT), orbitrap (electrostatic orbital trap), or the Fourier transform ion cyclotron resonance; but since they do not belong to equipments with acceptable prices, they are not widely used in common practice [35].

Costs efficiency of analytic procedures represent an important issue for all laboratories performing the residues analyses. The growth of capital expenses is conditioned and/or determined by: the use of systems enabling automation and shortening of analytic procedures; initial expensive equipment (TOF, Orbitrap, and others); testing a large number of samples per day; and other such factors. The alternative is to maximise the number of analytes identified by a single analytic procedure. This procedure, despite a high price, is definitely possible with the LC-MS/MS technique [72].

ULTRA PERFORMANCE LIQUID CHROMATOGRAPHY WITH TANDEM MASS SPECTROMETRY (UPLC-MS/MS)

The latest developments in the field of liquid chromatography are focused on: the development of new columns with a reduced size of the stationary phase; lower diameters; as well as, the use of higher pressures in the system. These developments significantly improve: the separation efficiency; shortens the duration of the analysis; improves the detection limits; reduces the consumption of solvents; and significantly reduces the elution times, which significantly increases the number of analysed samples per day. An example of this is the ultra performance liquid chromatography with tandem mass spectrometry (UPLC-MS/MS). This method is based on the principle of using the stationary phase consisting of particles smaller than 2.5 μm . Also well-known, is the combination of the UPLC and the TOF mass analyser for multi-residue analysis, providing the absolute and unequivocal confirmation of positive findings of coccidiostat residues in animal products [9], [36], [37], [74].

CONCLUSIONS

Laboratories performing certified inspection of residues must face the increasing number of stricter requirements and criteria applying to the performance of analytic methods as defined by the currently applicable legislation. The purpose of the legal requirements is to eliminate the risk resulting from the presence of residues in food matrices and provide the consumer with safe and full-value food. The highest priority is the development of accurate methods of residue identification in order to prevent residues entering the food chain and thus to protect the consumers' health.

History shows that the constant technical innovations and improvement of the performance of analytic methods used for the identification of coccidiostat residues in the food of animal origin and in feed, leads to the selection of the most suitable and appropriate methods. An example of

such advancement include the sensitive and reliable LC-MS/MS technique that identifies analytes on the basis of their molecule mass and structure. Nevertheless, it is necessary to realise that despite the significant improvements in this area, the extant analytic separation methods must be constantly improved.

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ASSESSMENT ON THE WELFARE AND ITS INFLUENCE ON THE HEALTH OF CART HORSES IN COMBOLCHA DISTRICT, ETHIOPIA

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ABSTRACT

A cross-sectional study on randomly selected 390 uncastrated cart horses was conducted from October 2011 to April 2012 to assess their major health and welfare problems in and around the town of Combolcha, Ethiopia. The welfare aspect of cart horses was determined in the study area by direct assessment, which included the incidence of lameness, injuries, and other diseases of cart horses in relationship to their body condition, age, and behaviour. Focus group discussions and proportional pulling techniques were also used to generate further information. The animals were allotted into three age groups; 0—6 years old (3.3%), 6—11 years old (49.4%), and 11—15 years old (46.9%) and their body condition was determined to be; thin (7.4%), fair (56.7%), good (32.6%), and fat (3.3%). The overall prevalence of injury was 32.1%. The prevalence of external injuries was found in animals; under 6 years old (61.5%), between 11—15 years of age (31.1%), and between 6 to 11 years of age (30.9%). In addition, more external injuries were registered in thin horses ($P < 0.05$) compared to horses having fair or good body condition. The least incidence of injury was recorded in fat horses. Significantly higher number of cases of injuries was reported in cart horses out of work, followed by cart horses transporting people and cart horses transporting goods (85.7%, 32.6%, and 21.2%, respectively; $X^2 = 12.142$; $df = 2$; $P < 0.05$). Further, an increased ($P < 0.01$) incidence of external injuries were observed in alert and not friendly horses. Improper harnesses, overworking, and overloading probably caused the injuries. The cart horses out of work were horses that could no

longer work due to severe injury and other serious health problems. They were abandoned by their owners and were straying in and around the town of Combolcha. The overall prevalence of lameness was 18.3%. Similar to the incidence of external injuries, the highest prevalence of lameness ($P = 0.780$) occurred in cart horses below 6 years old. In this study, animals with the fair body condition score, showed a high incidence of lameness (22.2%; $P = 0.075$). Further, cart horses out of work had a high incidence of lameness also (28.6%; $P = 0.461$), followed by cart horses used for goods transportation (23.1%) and cart horses used for people transportation (17.2%; $P = 0.795$). In addition, depressed horses had a high prevalence of lameness. Lameness found in this study most probably was caused by shoeing abnormalities in addition to the malconformation of the hoof. Despite the animals being affected by lameness, they were obliged to continue to work. In focus group discussions, respondents reported wounds as the primary health problem and the cause of the wounds as due to overloading of the cart horses. The study revealed that many cart horses working in the study area are experiencing multiple welfare problems.

Key words: age; animal welfare; behaviour; body condition; injury; lameness

INTRODUCTION

Horses are important draught animals for both the developed and developing countries of the world. They play an important role

in socio-economics by being used for farming, cart pulling, transportation, sporting and leisure activities. They are also important sources of food in certain countries. The American horse council estimated that horses have a direct impact on the economy of the United States of over \$39 billion and an indirect impact of over \$102 billion. Horses used to be, and still are, an important use of transportation and they do not pollute the earth [13].

According to a report compiled by the Food and Agricultural Organization of the United Nations [6], there are about 58 million horses in the world. Ethiopia is placed 8th in the world, and first in Africa, with its 1,655,385 horse population. Horses are efficient users of low quality, high fibre food and can tolerate up to 30% dehydration [21]. Compared to other production animals, they require lesser quantity of food and have the ability to conserve more energy. They drink infrequently and irregularly. This makes horses a suitable adaptable animal for harsh environments and difficult working conditions [17].

Horses provide a life line for rural and urban poor communities in Ethiopia. The report by the Brooke [1], an international charity, which works to improve the welfare of working equines around the world, indicated that horses are crucial to Ethiopian communities in rural and urban settings. This study revealed, working horses can reduce the burden for women while at the same time offering house holds an opportunity to generate income and improve food security by helping to reduce poverty [14]. According to Bifa and Woldemeskel [2], horses are important animals in the farming and transport systems of Ethiopia. As pack animals, they transport harvest from the fields to the homestead. They also transport agricultural products and fuel to the market and building materials and other commodities from the market to the villages. For rural traders and urban cart owners, horses are the sole means of income to sustain their extended families. The average net return from horse ownership and use has been estimated to be the equivalent of \$330 per year in Ethiopia [2]. Moreover, horses play a key role in job creation in the supply of; chains for carts, harnesses, health services, animal feed, hoof/shoeing, and other provisions.

The cart is a common form of short to mid distance transportation for people and goods in many small towns and cities in Ethiopia. A two wheel cart is pulled by one horse. Cart pulling horses are suffering from the burden of stress. While experiencing severe stress, they can succumb to disease [10]. It is known that animal health is a key component of animal welfare and the active involvement of veterinary services and veterinarians is important for improving animal welfare [12]. Despite cart horses' important socio-economic contributions, their welfare is often neglected and receive very low attention. Little interest has been given to research on welfare issues of the cart horses compared to donkeys in Ethiopia.

Therefore, the purpose of this study was to assess the welfare problems and their influence on the health of cart horses in and around Combolcha, Ethiopia.

MATERIALS AND METHODS

A cross sectional study was conducted on randomly selected 390 uncastrated cart horses residing in peri-urban and urban areas of the town of Combolcha, between October 2011 and April 2012. The sample size for this study was calculated based on the formula developed by Thrusfield [19].

$$n = \frac{1.96^2 \times p \times (1 - p)}{d^2}$$

Where n = sample size

p = expected prevalence = 50 % = 0.5

d = desired level of precision = 5 % = 0.05

$$n = \frac{1.96^2 \times 0.5 \times (1 - 0.5)}{0.05^2} = 384$$

Since there was no study previously carried out on cart horses and their welfare in the study area, 50% prevalence was used to calculate the sample size.

The welfare aspect of cart horses was determined in the study area by direct assessment, which included the incidence of lameness, injuries, and other diseases in association with body condition, age, and behaviour. The cart horses body condition was determined based on the method developed by Carroll and Huntington [4] as cited by Pritchard et al. [16]. The lameness was determined by observing their gait and the result was categorized as low grade, moderate, high grade and immobile to each lame cart horse based on the criteria formulated by Dyson [5]. The cart horses were physically examined for grossly visible skin/tissue injuries measuring greater than or equal to 5 cm located on any part of the animal body. Injuries had to be active, with ongoing tissue damage, with or without blood/exudates/pus, abscess formation, or any secondary bacterial complication as stated by Biffa and Woldemeskel [2]. The age of the animals was estimated by birth records obtained from the owners and dentition characteristics. Accordingly, the cart horses were allotted into four age groups: Group 1 consisted of animals 0–6 years old; Group 2, 6–11 year old; Group 3, 11–15 years old; and Group 4, > 15 years old. The behaviour of the study animals was investigated by close observation and contact with the animals, both at work and during rest. Their behaviour was characterized as; anxious, friendly, alert and friendly, alert and not friendly, or depressed [2]. Other diseases manifested by the study of the cart horses were diagnosed in the veterinary clinic and recorded as well.

Focus group discussions were held with local communities, cart horse owners, and clients for the indirect assessment of the welfare of the cart horses. The discussions were focused on: the role of the cart horses, management constraints, cart horses health problems, handling of cart horses by cartmen while working and the reaction of the community, government or non-governmental organizations (NGOs) to the mishandling of the cart horses. Information gathered from the focus group discussions were given a rank.

The data collected from the study area was entered into MS excel spread sheets and analysed using STATA 7.0 and SPSS 15.0 statistical software. The statistical analysis used included the comparison of proportions, an X^2 test and a test of agreement [19]. The Chi square test was applied to test if any statistical association existed between the welfare of the cart horses and the risk factors, such as; age, body condition, and type of work the horses performed.

RESULTS

During the study period, the animals were clinically examined for the presence of any gross external lesions, lameness, and other diseases. The body condition of the animals was scored as well as, their age, and their behaviour. All of the study cart horses were male. Of the study cart horses, 3.3 % were under the age of 6 years, whereas 49.4 % were between 6—11 years and 46.9 % between 11—15 years of age. There were no cart horses above 15 years of age (Table 1.).

Table 1. Age grouping of cart horses

| Age [year] | Number of animals | Percentage [%] |
|------------|-------------------|----------------|
| 0—6 | 13 | 3.3 |
| 6—11 | 194 | 49.4 |
| 11—15 | 183 | 46.9 |
| > 15 | 0 | 0 |

A study of the type of work revealed that 84.4 % of the horses were used for people transport, 13.3 % for transport of goods, and 1.8 % were found to be out of work (Table 2).

Table 2. Type of work the study animals were engaged in

| Type of work | Number of animals | Percentage [%] |
|---------------------|-------------------|----------------|
| Transporting people | 331 | 84.4 |
| Transporting goods | 52 | 13.3 |
| Out of work | 7 | 1.8 |

Of the 390 cart horses examined, the body conditions of 7.4 %, 56.7 %, 32.6 %, and 3.3 % were found to be thin, fair, good, and fat, respectively (Table 3.).

Table 3. Body condition score of the study animals

| Body condition score | Number of animals | Percentage [%] |
|----------------------|-------------------|----------------|
| Thin | 29 | 7.4 |
| Fair | 221 | 56.7 |
| Good | 127 | 32.6 |
| Fat | 13 | 3.3 |
| Very fat | 0 | 0 |

The highest number of horses (66.9 %) demonstrated alert and not friendly behaviour followed by depressed (14.9 %), anxious (8.7 %), friendly (8.2 %), and alert and friendly behaviour (1.3 %) (Table 4.).

Table 4. Observed behaviour of the study animals

| Behaviour | Number of animals | Percentage [%] |
|------------------------|-------------------|----------------|
| Friendly | 32 | 8.2 |
| Alert and friendly | 5 | 1.3 |
| Alert and not friendly | 261 | 66.9 |
| Depressed | 58 | 14.9 |
| Anxious | 34 | 8.7 |

Of the cart horses, a total of 32.1 % (n = 125) had injuries on different parts of their bodies (Fig. 1).

Injuries on the cart horses were more frequently observed on girth (12.8 %; n = 50), shoulder (3.30 %; n = 16), and mouth/bit (4.9 %; n = 19) compared with other body parts (Fig. 1). The association of the prevalence of external injuries on cart horses with age, body condition, work type, and behaviour is illustrated in Table 5. An increased prevalence of external injuries was found in animals under 6 years old (61.5 %) compared to animals between 11—16 years of age (31.1 %) and 6—11 years old (30.9 %) in a descending order. According to this study, there was no significant difference in the prevalence of external injuries among different age groups of animals ($X^2 = 5.371$, $df = 3$, $p = 0.068$).

More external injuries were registered in thin horses compared to horses having fair or good body condition (58.6 %, 33.9 %, and 24.4 %, respectively). The least incidence of injuries was recorded in fat horses (15.4 %). The body condition was found to significantly influence the prevalence of external injuries ($X^2 = 14.824$, $df = 4$, $P < 0.05$). High cases of injuries were reported on cart horses out of work followed by cart horses transporting people and cart horses transporting goods (85.7 %, 32.6 %, and 21.2 %, respectively). In the present study, significant variation in the prevalence of external injury on cart horses performing different type of work was encountered ($X^2 = 12.142$, $df = 2$, $P < 0.05$). Significantly higher ($P < 0.01$) incidence of external injuries was observed in alert and not friendly horses comparing to depressed, friendly, and anxious horses (100 %, 51.7 %, 40.6 %, and 38.2 %, respectively). The least prevalence of injuries was recorded in alert and friendly cart horses (24.5 %). The behaviour of the cart horses was found to be highly associated with the prevalence of injury ($X^2 = 29.392$).

The overall prevalence of lameness among the study animals (n = 390) was 18.3 % (n = 71). From the affected animals, 36.6 % had high grade, 28.2 % moderate, and 33.8 % low grade lameness, while 1.4 % of them were immobile (Table 6.).

The association of the prevalence of lameness in cart horses with age, body condition, work type and behaviour is indicated in Table 7.

The lowest prevalence of lameness occurred in cart horses below 6 years of age (3.0 %; n = 13) followed by, 6—11 years old (33.0 %; n = 194), and 11—15 years old (35.0 %; n = 183). There was no significant difference among different age groups in the prevalence of lameness. On the other hand,

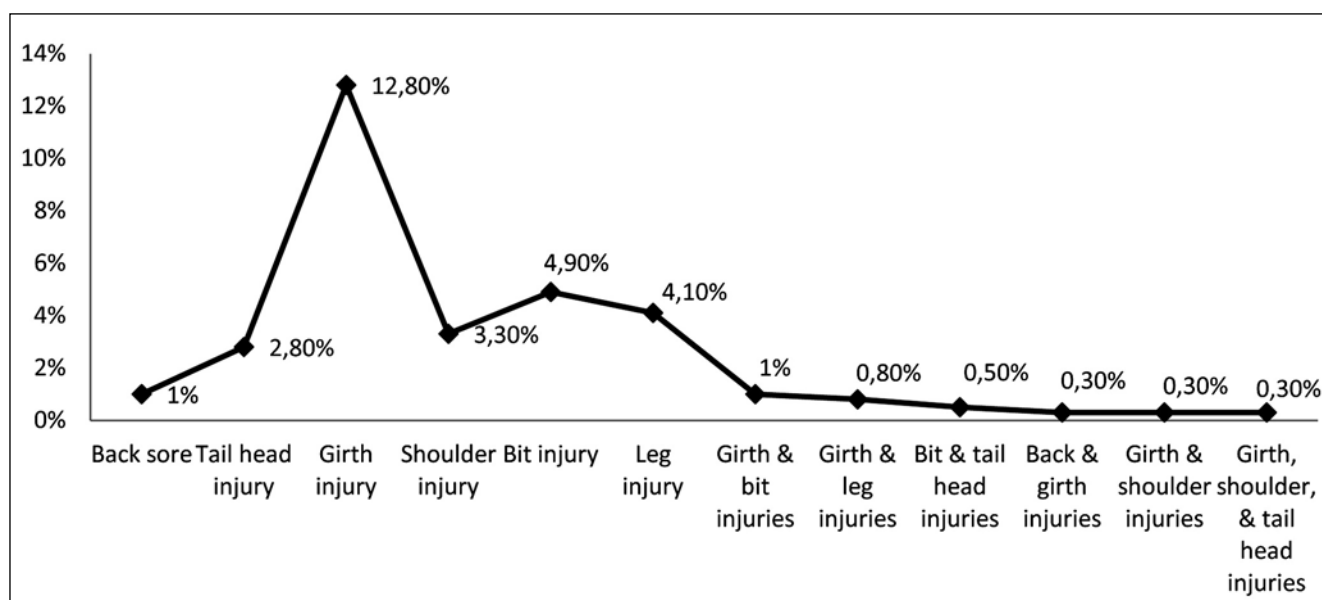


Fig. 1. Prevalence of injuries on different body parts of cart horses

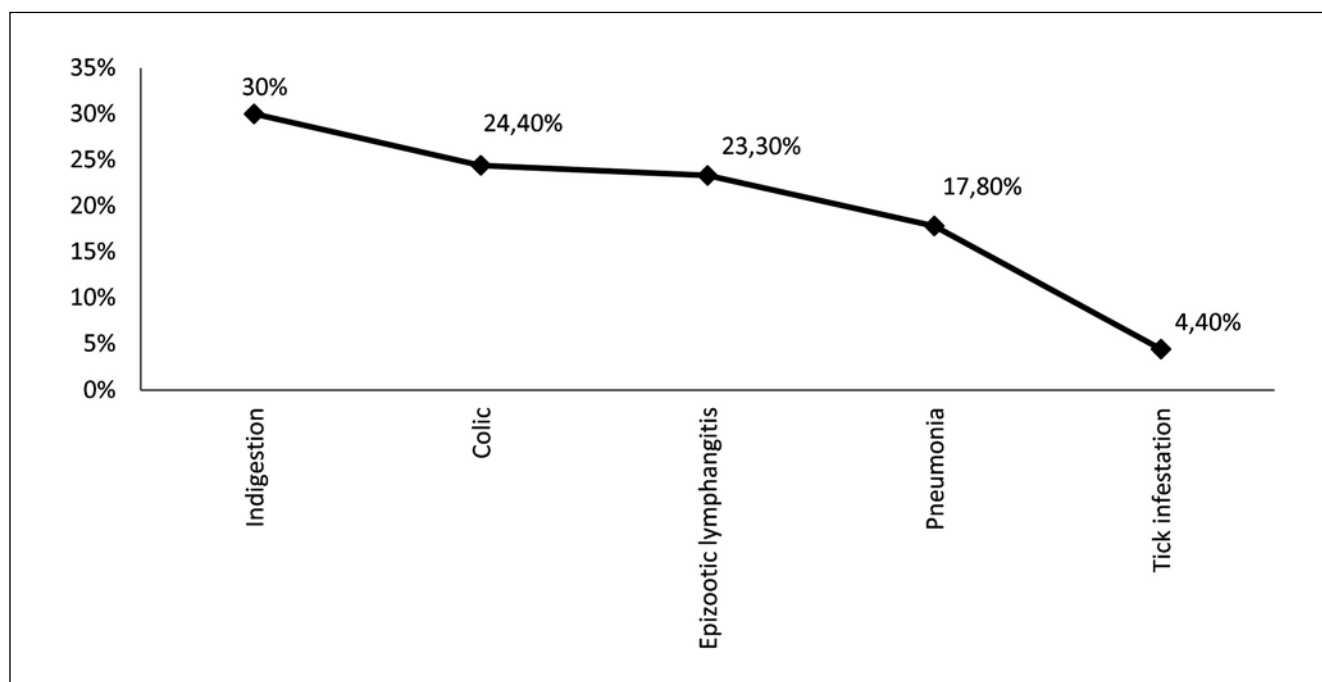


Fig. 2. Prevalence of other diseases in the cart horses

animals with fair body condition score showed high incidence of lameness (22.2 %) compared to thin (20.7 %), good (11.8 %), and fat (7.7 %) in decreasing order. The Chi-square value of 6.907 indicated an association of body condition and lameness. However, the prevalence of lameness did not differ significantly among animals with different body condition score.

Cart horses out of work had a high incidence of lameness (28.6 %), followed by cart horses used for goods trans-

portation (23.1 %) and cart horses used for people transportation (17.2 %). The alert and not friendly animals had a high prevalence of lameness (24.1 %), followed by anxious (20.6 %), alert and friendly (20 %), depressed (17.2 %), and friendly (12.5 %). The incidence of lameness by behaviour and work type was not significant.

A total of 90 (23.1 %) sick study cart horses were examined in Combolcha town veterinary clinic. Of which, 27 (30 %) were affected with indigestion, 22 (24.4 %) with colic,

Table 5. Prevalence of external injuries by age, body conditions (BCS), type of work, and behaviour

| Factor | Factor categories | Prevalence [%] | X ² = value | df | P value |
|------------|--------------------------------|----------------|------------------------|----|---------|
| Age (year) | 0—6 (n = 13) | 61.5 | 5.371 | 2 | 0.068 |
| | 6—11 (n = 194) | 30.9 | | | |
| | 11—15 (n = 183) | 31.1 | | | |
| BCS | Thin (n = 29) | 58.6 | 14.824 | 3 | 0.02 |
| | Fair (n = 221) | 33.9 | | | |
| | Good (n = 127) | 24.4 | | | |
| | Fat (n = 13) | 15.4 | | | |
| Work type | Transport people (n = 331) | 32.6 | 12.142 | 2 | 0.002 |
| | Transport goods (n = 52) | 21.2 | | | |
| | Out of work (n = 7) | 85.7 | | | |
| Behaviour | Friendly (n = 32) | 40.6 | 29.392 | 4 | 0.00 |
| | Alert and friendly (n = 261) | 24.5 | | | |
| | Alert and not friendly (n = 5) | 100 | | | |
| | Anxious (n = 34) | 38.2 | | | |
| | Depressed (n = 58) | 51.7 | | | |

df = degree of freedom; X² = Chi square; significant at P < 0.05

Table 6. Prevalence of lameness in the study animals

| Degree of lameness | Number of animals | Prevalence in % |
|--------------------|-------------------|-----------------|
| High | 26 | 36.6 |
| Moderate | 20 | 28.2 |
| Low | 24 | 33.8 |
| Immobile | 1 | 1.4 |

21 (23.3 %) with epizootic lymphangitis, 16 (17.8 %) with pneumonia, and 4 (4.4 %) with tick infestation (Fig. 2).

In the focus group discussions, it was found that inadequate feed supply to cart horses was a major management constraint (15.3 % of respondents), followed by poor veterinary health service (14.5 % of respondents), and inappropriate harnessing (12.7 % of respondents). Respondents further reported wound as the primary health problem of the cart horses (16.8 % of respondents) and cause of the wounds were overloading cart horses (11.1 % of the respondents) followed by the beating of the cart horses by the cart drivers (10.7 % of respondents).

DISCUSSION

Cart horses are the most important animals for transportation of people and their goods both in urban and rural areas of developing countries [7]. In the peri-urban and urban areas of Ethiopia, horses mostly pull carts. They are important and the sole means of livelihood for many families. They work in unfriendly environmental conditions, including intense heat, difficult terrain, and inappropriate harnesses [9]. Despite their important socio-economic contributions, cart horses are often treated harshly; suffer from overloading and overworking coupled with poor shelter, malnutrition, and poor health condition.

In this investigation, 32.1 % (n = 125) of the study cart horses (n = 390) had incidences of external injuries on different part of their bodies. The incidence of injuries in this study was lower than the prevalence of injuries indicated by Biffa and Woldemeskel [2]. Injuries in the study cart horses were more frequently observed on the girth (12.8 %), shoulder (4.9 %), and mouth/bit (4.1 %), compared with other body parts. The injuries on the girths were probably caused by improper harnesses. Harnesses of carts in the study area were mostly made up of strips of old car tyres. Despite their endurance, strips of car tyres were inflexible and had sharp edges. Thus, due to their constant friction with the girth of the animals, they might cut into the skin and caused injury. The shoulder injuries were probably caused by iron or

Table 7. Prevalence of lameness in association with age, body condition, type of work and behaviour of study cart horses

| Factors | Factor categories [n] | Prevalence [%] | X ² -value | df | p-value |
|-------------------|----------------------------|----------------|-----------------------|----|---------|
| Age (year) | 0—6 (13) | 3.0 | 0.497 | 2 | 0.780 |
| | 6—11 (194) | 33.0 | | | |
| | 11—15 (183) | 35.0 | | | |
| BCS | Thin (29) | 20.7 | 6.907 | 3 | 0.075 |
| | Fair (221) | 22.2 | | | |
| | Good (127) | 11.8 | | | |
| | Fat (13) | 7.7 | | | |
| Work type | Transport people (331) | 17.2 | 1.549 | 2 | 0.461 |
| | Transport goods (52) | 23.1 | | | |
| | Out of work (7) | 28.6 | | | |
| Behaviour | Friendly (32) | 12.5 | 2.378 | 4 | 0.795 |
| | Alert and friendly (261) | 20.0 | | | |
| | Alert and not friendly (5) | 24.1 | | | |
| | Anxious (34) | 20.6 | | | |
| | Depressed (58) | 17.2 | | | |

wood made saddles firmly tied to the body without adequate protection underneath, coupled with overloading and overworking. The saddle might constantly rub and irritated the shoulder of the animal causing injury. The cart horses usually transported four people being overloaded in a trip and worked for 6 to 7 hours long a day. In agreement with our study, Yilma et al. [20] and Biffa and Woldemeskel [2] encountered injuries in equines caused by improper harnesses and saddles. Pearson [15] reported that overweight and different types of load/work contributed to high causes of back sores in the donkeys of central Ethiopia. Fred and Pascal [8] also reported that donkeys in Kenya developed extensive sores and wounds due to overworking.

Contrary to the findings of Biffa and Woldemeskel [2], higher prevalences of injuries were registered in horses less than 6 years of age compared to older horses. Significantly ($P < 0.05$) more external injuries were registered in thin horses compared to horses having fair or good general body conditions. The least incidence of injury was recorded in fat horses. This may be because thin horses are not tolerant to hardship conditions and injuries persist longer due to weak immune reactions. In addition, where the horse is in poor body condition and lacks the layer of subcutaneous fat, there will be a greater prevalence of sores due to ill fitting or badly made harnesses [3], [9]. In addition, significantly higher ($P < 0.05$) cases of injuries were reported in cart horses out of work, followed by cart horses transporting people

and cart horses transporting goods, in decreasing order. The cart horses out of work were horses that can no longer work due to severe injury and other severe health problems. They were abandoned by their owners and were straying on the roads in and around Combolcha. Also, significantly higher ($P < 0.01$) incidence of external injuries was observed in alert and not friendly horses, compared to depressed, friendly, and anxious horses. Because of working under difficult conditions, cart horses were unable to express natural behaviour. When they become aggressive, they may inflict injury to themselves. Forceful and reckless usage of the bit to lead and brake cart horses probably caused injuries at the commissures of the mouth. Nawaz et al. [11] supported this finding.

Similar to the prevalence of injuries, a high incidence of lameness was recorded in young animals less than six years of age. Thin horses showed an insignificantly higher prevalence of lameness followed by horses that had fair body condition. Further, depressed animals had a high prevalence of lameness compared to anxious, alert and not friendly, alert and friendly, and friendly cart horses in descending order. In the study area, cart horses were shod with rubber taken from old car tyre. The hoof/shoeing was performed by low skilled farriers who did not understand hoof balance and care. Therefore, lameness found in this study most probably was caused by hoof/shoeing abnormalities in addition to the possible malconformation of a hoof. This finding was sup-

ported by the study of Tadich [18]. Despite the animals being affected by lameness, they were obliged to continue to work. Because these animals were not moving fast to the desire of the cart drivers, they were often whipped and exposed to further injury and severe stress. The lame animals in this study were obliged to work because if they stopped working, the cart owners and their families might lose their livelihood.

The inadequate provision of water and feed to cart horses was manifested by low body condition scores. Sixty four point one percent (64.1 %) of the study animals scored thin to fair body condition. In addition, indicators of poor clinical health of the study animals were recorded on visits to the veterinary clinic including; inappetence, colic, pneumonia, lymphangitis, and parasitic infestations.

Respondents from the focus group discussions gave importance to the provision of adequate feed and water to the cart horses, usage of appropriate harnesses and carts, provision of adequate veterinary service, and avoidance of whipping and overworking to improve the welfare of cart horses.

CONCLUSIONS

The lack of knowledge of animal welfare and the outcome of poor welfare and the poverty of cart horse's owners leads to multiple welfare and health problems of cart horses. Cart horses were routinely exposed to stressful situations by; overworking, overloading, working in hot weather, and difficult terrain without adequate water and feed supply. Further, the welfare and health of most cart horses was significantly compromised by the injuries caused by improper harnesses and hoof/shoeing. Direct physical abuse of the cart horses was often observed. Even though the horses were lame and/or sick with other diseases, they were obliged to continue to work without any treatment. Moreover, when the cart horses were deemed unable to work because of disease and exhaustion, they were abandoned by their owners and left to stray on the roads in the suburbs of Combolcha town.

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EFFECT OF ENROFLOXACIN ON ACEPROMAZINE-KETAMINE ANAESTHESIA IN RABBITS

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ABSTRACT

Time to induction, duration of recumbency and time to standing, as well as changes in heart rate (HR), respiratory frequency (f_R) and rectal temperature (RT) over a period of 120 min were determined in 12 healthy rabbits, after intramuscular (IM) injection of 5 mg.kg⁻¹ acepromazine, followed 30 min later with concurrent IM injections of 75 mg.kg⁻¹ ketamine and the antibiotic, 10 mg.kg⁻¹ enrofloxacin (AKE). Two weeks later, the rabbits were similarly injected with acepromazine and ketamine (AK) but without enrofloxacin to serve as a control. The time to induction (1.8 ± 0.3 min) with AKE rabbits was significantly ($P < 0.05$) shorter than that with the AK value of 4.4 ± 0.8 min, while the duration of recumbency (149.0 ± 5.6 min) and time to standing (20.2 ± 6.5 min) with AKE rabbits were significantly ($P < 0.05$) longer than their respective AK values of 139.4 ± 4.9 min and 13.6 ± 2.0 min. With the AKE, the mean HR ranged from 164.0 ± 7.2 to 192.7 ± 16.1 beats.min⁻¹ as against the control range of 176.6 ± 4.2 to 206.6 ± 5.6 beats.min⁻¹; the mean f_R ranged from 26.7 ± 3.2 to 56.7 ± 10.7 breaths.min⁻¹ as against the control range of 28.6 ± 7.9 to 37.0 ± 4.1 breaths.min⁻¹; and the mean RT ranged from 38.1 ± 0.3 to 39.6 ± 0.6 °C as against the control range of 39.3 ± 0.2 to 39.7 ± 0.2 °C. In all treatment groups, the recorded changes in physiological parameters fell within the normal range of values for awake rabbits. It was concluded that the concurrent administration of enrofloxacin and acepromazine-ketamine anaesthesia in clinically healthy rabbits produced shorter time to induction, but longer duration of recumbency and time to standing

than the corresponding control values. Significantly lower mean heart rate and rectal temperature as well as higher respiratory frequency were also recorded compared to the control values.

Key words: acepromazine; anaesthesia; concurrent; enrofloxacin; ketamine; rabbits

INTRODUCTION

Ketamine is frequently used to provide chemical restraint in animals [17], [18], [29], [42]. It is a useful injectable anaesthetic agent because of its: wide margin of safety; compatibility with most other anaesthetic agents; the possibility of administering it by the intramuscular route where venepuncture proves difficult to achieve in the awake animal; and its sympathomimetic properties causing an increase in heart rate, cardiac output and blood pressure [18], [20], [29]. However, when used alone, ketamine produces poor muscle relaxation, hypertonus, persistent pain reflex responses, and violent recovery from anaesthesia [17], [18], [20], [29], [42]. In order to counteract these side effects of ketamine, sedatives including an α_2 adrenergic agonist (xylazine or medetomidine), acepromazine or a benzodiazepine (diazepam, midazolam, or zolazepam) have been used in combination with it [3], [29], [32]. Indeed, ketamine-based combinations are currently the most popular in rabbit anaesthesia [3], [11], [24], [34].

Enrofloxacin is a synthetic chemotherapeutic agent of the fluoroquinolone family. The drug has a broad-spectrum antibacterial

activity and it has been licensed for use in the treatment of bacterial infections in the rabbit [22], [36]. Therefore, in difficult practice situations where perioperative bacterial culture and sensitivity testing is not practical, enrofloxacin may be administered to rabbits for its broad-spectrum antibacterial activity to prevent or treat infections. Occasionally, rabbits on current enrofloxacin medication may be scheduled for clinical procedures necessitating the use of anaesthetics, thus providing the opportunity for an antimicrobial-anaesthetic interaction to occur. Such drug-drug interaction could lead to alteration in the actions or frank toxicity of the drugs involved [9], [33]. In one study, in cats pretreated with chloramphenicol, A d e t u n j i et al. [1] reported that xylazine-ketamine anaesthesia was associated with longer duration of analgesia, sleeping and standing times. Also in rabbits with concurrent administration of enrofloxacin and xylazine-ketamine anaesthesia, A d e t u n j i and L a w a l [2] reported shorter times to induction and standing, but a longer duration of recumbency. However, the effect of concurrent administration of enrofloxacin on acepromazine-ketamine anaesthesia has yet to be reported in the rabbit.

The aim of this study, therefore, was to determine the effects, if any, of the concurrent administration of enrofloxacin (E) on acepromazine-ketamine (AK) anaesthesia in rabbits, using selected anaesthetic indices, as well as heart rate (HR), respiratory frequency (f_R) and rectal temperature (RT) responses, to assess the degree of depression of both life support systems, in the course of the trials.

MATERIALS AND METHODS

Animals

Twelve adult New Zealand white and American Chinchilla cross rabbits (six bucks and six does) whose body weight ranged from 1.1 to 1.5 kg (1.3 ± 0.1 kg, mean \pm SD), were used for this study. They were obtained from the experimental animal unit of our Faculty of Veterinary Medicine. The animals were maintained in individual cages with dimensions of $60 \times 60 \times 60$ cm, for at least four weeks before the experiments for acclimation and constant human handling. They were fed *ad libitum* with commercial pellet food and fresh water was provided free choice in their cages.

Study design

The Institutional Animal Care and Use Committee of our Faculty approved this study. Two series of experiments were carried out on the 12 experimental rabbits at two weeks intervals. The first series being the treatment consisted of the intramuscular (IM) administration of acepromazine, ketamine and enrofloxacin (AKE), while the second series (control) were similarly treated but without the enrofloxacin injection (AK). Selected anaesthetic indices were calculated for both groups. Some physiological parameters of the rabbits were also measured immediately before the concurrent administration of ketamine and enrofloxacin, and subsequently at 10 min interval until the effect of the drugs abated.

Experimental procedure

Food and water were allowed up to the time of the trials. Each rabbit was weighed using a special weighing balance with a reading range of 0.0 to 5.0 kg. Each rabbit in the AKE series was premedicated with IM injection of 5 mg.kg^{-1} of acepromazine (Berkuce^(R)

10 mg.ml^{-1} , Berk Pharmaceutical Ltd, England), followed 30 min later by concurrent IM injections of 75 mg.kg^{-1} of ketamine (Ketmin^(R) 50 mg.ml^{-1} , Laborate Pharmaceutical, India) and 10 mg.kg^{-1} of enrofloxacin (Conflex^(R) Vet 50 mg.ml^{-1} , Concept Pharmaceutical Ltd, India) into either thigh [13], [22].

Rabbits in the control series (AK) were similarly treated but without the enrofloxacin injection. The anaesthetized rabbits were placed on their right lateral recumbency on a foam padded wooden table.

Calculated anaesthetic indices

In these trials, the following selected anaesthetic indices were calculated:

- 1. Time to induction:** time interval (in min) between the IM injection of ketamine and the loss of righting reflex by the rabbit.
- 2. Duration of recumbency:** time interval (in min) between the loss of righting reflex and assumption of sternal posture by the rabbit.
- 3. Time to standing:** time interval (in min) between the assumption of sternal and standing postures by the rabbit.

Measured physiological parameters

The baseline HR, f_R , and RT were measured immediately before concurrent administration of ketamine and enrofloxacin (time 0 min), and subsequently at 10 min interval until the effects of the drugs abated. The HR in beats.min^{-1} was evaluated with the aid of a precordial stethoscope. The f_R in breaths.min^{-1} was determined by counting the costo-abdominal movements of the anaesthetized rabbits. The RT ($^{\circ}\text{C}$) was measured using a digital thermometer inserted into the rectum.

Statistical analysis

The data were expressed as the mean \pm SD of 12 rabbits. The mean values of the selected anaesthetic indices of both the AKE and AK groups were compared using the student's t-test for paired data. The means of measured physiological parameters of the AKE were also compared with the AK values using analysis of variance (ANOVA) for repeated measures followed by Tukey multiple comparisons test as post-test. A value of $P < 0.05$ was accepted as statistically significant. The NCSS 2004 statistical package was used (10).

RESULTS

Calculated anaesthetic indices

The time to induction (1.8 ± 0.3 min) with AKE rabbits was significantly ($P < 0.05$) shorter than with AK value of 4.4 ± 0.8 min, while the duration of recumbency (149.0 ± 5.6 min) and time to standing (20.2 ± 6.5 min) with AKE rabbits were significantly ($P < 0.05$) longer than their respective AK values of 139.4 ± 4.9 min and 13.6 ± 2.0 min.

Measured physiological parameters

The mean HR, f_R , and RT for the AKE and AK series were as shown in Tables 1, 2 and 3 respectively. The mean HR with the AKE ranged from 164.0 ± 7.2 to $192.7 \pm 16.1 \text{ beats.min}^{-1}$ as against the control range of 176.6 ± 4.2 to $206.6 \pm 5.6 \text{ beats.min}^{-1}$. Apart from the 50th min time interval, the AKE rabbits had

Table 1. Heart rate responses of rabbits to intramuscular administration of either acepromazine-ketamine (AK) or acepromazine-ketamine-enrofloxacin (AKE)

| Time interval [min] | HR [beats.min ⁻¹] | |
|---------------------|-------------------------------|---------------------------|
| | AK | AKE |
| 0 ^a | 198.2 ± 4.4 | 192.7 ± 16.1 |
| 10 | 206.6 ± 5.6 ⁺ | 185.3 ± 19.4 [*] |
| 20 | 189.5 ± 2.5 ⁺ | 188.7 ± 13.7 |
| 30 | 189.2 ± 7.3 ⁺ | 184.0 ± 6.1 [*] |
| 40 | 180.0 ± 3.9 ⁺ | 177.3 ± 6.6 ⁺ |
| 50 | 176.6 ± 4.2 ⁺ | 180.0 ± 3.3 [*] |
| 60 | 186.1 ± 4.8 ⁺ | 172.7 ± 6.2 ^{*+} |
| 70 | 188.3 ± 6.4 ⁺ | 180.7 ± 8.9 ^{1*} |
| 80 | 186.1 ± 5.9 ⁺ | 178.0 ± 7.4 [*] |
| 90 | 190.5 ± 3.4 ⁺ | 182.0 ± 9.6 [*] |
| 100 | 199.3 ± 3.7 | 182.0 ± 9.6 [*] |
| 110 | 200.0 ± 7.4 | 176.0 ± 8.5 ⁺ |
| 120 | 200.0 ± 6.2 | 164.0 ± 7.2 ^{*+} |

Data are expressed as mean ± SD of 12 rabbits

AK: IM injection of 5 mg.kg⁻¹ acepromazine, followed 30 min later by IM injection of 75 mg.kg⁻¹ ketamine; **AKE:** IM injection of 5 mg.kg⁻¹ acepromazine, followed 30 min later by concurrent IM injections of 75 mg.kg⁻¹ ketamine and 10 mg.kg⁻¹ enrofloxacin

^a — Baseline data obtained immediately before concurrent administration of ketamine and enrofloxacin;

^{*} — P < 0.05 versus control;

⁺ — P < 0.05 versus baseline data

significantly ($P < 0.05$) lower mean HR than the corresponding control values throughout the trial period. Whereas the mean HR for AKE fell significantly from the baseline value at 40, 60, 110 and 120 min time intervals, the mean HR for AK fell significantly in the first 90 min of the trial and returns to the baseline value after 100th min time interval (Table 1). The mean f_R with the AKE ranged from 26.7 ± 3.2 to 56.7 ± 10.7 breaths.min⁻¹ as against the control range of 28.6 ± 7.9 to 37.0 ± 4.1 breaths.min⁻¹. Although they maintained similar mean f_R values up to the 30th min time interval, thereafter, AKE rabbits had significantly higher mean f_R than the corresponding control values until the end of the trial period. In addition, the mean f_R with the AKE increased significantly more than the baseline value from the 40th min time interval until the end of the trial, while the AK rabbits maintained similar mean f_R values throughout the trial period (Table 2). The mean RT with the AKE ranged

Table 2. Respiratory frequency responses of rabbits to intramuscular administration of either acepromazine-ketamine (AK) or acepromazine-ketamine-enrofloxacin (AKE)

| Time interval (min) | f_R [beats.min ⁻¹] | |
|---------------------|----------------------------------|--------------------------|
| | AK | AKE |
| 0 ^a | 32.0 ± 10.4 | 30.3 ± 2.9 |
| 10 | 28.8 ± 4.5 | 26.7 ± 3.2 |
| 20 | 29.7 ± 5.9 | 28.7 ± 4.6 |
| 30 | 30.6 ± 8.5 | 32.3 ± 4.0 |
| 40 | 28.6 ± 7.9 | 38.3 ± 4.1 [*] |
| 50 | 31.1 ± 3.6 | 48.0 ± 5.8 ⁺ |
| 60 | 35.2 ± 4.6 | 50.0 ± 6.8 ⁺ |
| 70 | 32.7 ± 2.7 | 56.7 ± 10.7 ⁺ |
| 80 | 37.0 ± 4.1 | 46.0 ± 7.0 ⁺ |
| 90 | 33.1 ± 2.7 | 49.3 ± 5.9 ⁺ |
| 100 | 35.3 ± 3.1 | 53.0 ± 10.9 ⁺ |
| 110 | 35.7 ± 4.6 | 48.7 ± 7.7 ⁺ |
| 120 | 35.2 ± 3.9 | 44.7 ± 5.0 ⁺ |

Data are expressed as mean ± SD of 12 rabbits

AK: IM injection of 5 mg.kg⁻¹ acepromazine, followed 30 min later by IM injection of 75 mg.kg⁻¹ ketamine; **AKE:** IM injection of 5 mg.kg⁻¹ acepromazine, followed 30 min later by concurrent IM injections of 75 mg.kg⁻¹ ketamine and 10 mg.kg⁻¹ enrofloxacin

^a — Baseline data obtained immediately before concurrent administration of ketamine and enrofloxacin; ^{*} — P < 0.05 versus control;

⁺ — P < 0.05 versus baseline data

from 38.1 ± 0.3 to 39.6 ± 0.6 °C as against the control range of 39.3 ± 0.2 to 39.7 ± 0.2 °C. Conversely, at the 20 min time interval, the mean RT of the AKE rabbits fell significantly from the baseline value and between the control values until the end of the trial, whereas AK rabbits maintained their mean RT throughout the trial period, except at the 40th min time interval (Table 3). However, in both groups, recorded changes in physiological parameters fell within the normal ranges of HR (130—325 beats.min⁻¹), f_R (32—60 breaths.min⁻¹), and RT (38.5—40.0 °C) for awake rabbits (21, 23).

DISCUSSION

The results of this study showed that concurrent administration of enrofloxacin and acepromazine-ketamine anaesthesia in clinically healthy rabbits produced shorter time to

Table 3. Rectal temperature responses of rabbits to intramuscular administration of either acepromazine-ketamine (AK) or acepromazine-ketamine-enrofloxacin (AKE)

| Time interval [min] | RT [°C] | |
|------------------------|-------------------------|-------------------------|
| | AK | AKE |
| 0 ^a | 39.3 ± 0.2 | 39.6 ± 0.6 |
| 10 | 39.4 ± 0.2 | 39.3 ± 0.4 |
| 20 | 39.5 ± 0.2 | 39.0 ± 0.4 ⁺ |
| 30 | 39.5 ± 0.2 | 39.0 ± 0.3 ⁺ |
| 40 | 39.7 ± 0.2 ⁺ | 38.9 ± 0.4 ⁺ |
| 50 | 39.6 ± 0.2 | 38.9 ± 0.3 ⁺ |
| 60 | 39.6 ± 0.2 | 38.9 ± 0.4 ⁺ |
| 70 | 39.6 ± 0.2 | 38.8 ± 0.5 ⁺ |
| 80 | 39.4 ± 0.4 | 38.7 ± 0.4 ⁺ |
| 90 | 39.4 ± 0.1 | 38.2 ± 0.4 ⁺ |
| 100 | 39.4 ± 0.2 | 38.1 ± 0.3 ⁺ |
| 110 | 39.5 ± 0.2 | 38.4 ± 0.4 ⁺ |
| 120 | 39.3 ± 0.2 | 38.2 ± 0.4 ⁺ |

Data are expressed as mean ± SD of 12 rabbits

AK: IM injection of 5 mg.kg⁻¹ acepromazine, followed 30 min later by IM injection of 75 mg.kg⁻¹ ketamine; **AKE:** IM injection of 5 mg.kg⁻¹ acepromazine, followed 30 min later by concurrent IM injections of 75 mg.kg⁻¹ ketamine and 1 mg.kg⁻¹ enrofloxacin

^a — Baseline data obtained immediately before concurrent administration of ketamine and enrofloxacin;

⁺ — P < 0.05 versus control; ⁺ — P < 0.05 versus baseline data

induction, but longer duration of recumbency and time to standing; with significant differences in heart rate, respiratory frequency and rectal temperature when compared with the control values.

The experimental rabbits in this study were not deprived of food and water before anaesthetic induction as recommended by Flecknell [14]. This is because rabbits are not known to vomit during either induction of, or recovery from, anaesthesia, so that there is no risk of aspiration by the anaesthetized rabbits. Furthermore, withholding food for any length of time would predispose the rabbit to the development of hypoglycaemia or trigger digestive disturbances that could result in endotoxaemia [14]. Also, deprivation of water from rabbits is known to cause preanaesthetic dehydration that is often difficult to control [14]. The intramuscular doses of acepromazine (5 mg.kg⁻¹) and ketamine (75 mg.kg⁻¹) employed in this study were those recommended by Flecknell [13] for clinical use in rabbits and the dosing interval of

30 min were those recommended by Clifford [7] for rabbits. It is recognised that drug doses are preferably tailored to the need of the patient in real clinical practice. For the same reason, a fixed dose of 10 mg.kg⁻¹ of enrofloxacin [22], [36] was employed in this study. Although the subcutaneous route of administration of the antimicrobial agent was recommended by these authors in the rabbit, the intramuscular route was preferred in this study for prophylaxis at the time of anaesthesia in order to achieve high blood level of enrofloxacin at the time of surgery.

The anaesthetized rabbits were not tracheally intubated because unlike in dogs, it is difficult to establish in rabbits, due to their long and narrow oral cavity [18], [21]. Furthermore, this mimics the current clinical practice where endotracheal intubation is not routinely carried out in rabbits [15]. The fact that no respiratory complications were recorded in the course of the trials suggests that endotracheal intubation is not crucial to maintaining an open airway in the anaesthetized rabbits.

In this study, the finding that the mean time to induction of the AKE-treated rabbits was shorter than the control value indicates a more rapid uptake of drugs from the IM site of injection. The mechanism by which this was achieved is not clear. Enrofloxacin is known to be rapidly absorbed from both oral and parenteral sites of administration [36]. It is probable that the antimicrobial agent crosses the blood-brain barrier in small amounts to cause some depression of the central nervous system, leading to the rapid onset of anaesthesia in the anaesthetized rabbits. This is likely to be true, since 6-10 percent of enrofloxacin present in the serum is reportedly found in the cerebrospinal fluid [36].

This difference in the induction times between the treatment and control groups, however, should not be of clinical concern. The longer duration of recumbency and time to standing associated with AKE-treated rabbits than the control values, suggested that the antimicrobial agent has some potentiating effects on acepromazine-ketamine anaesthesia in the rabbit. In dogs and cats, ketamine and enrofloxacin are both known to be renally excreted in part and hepatically metabolised [16], [18], [29], [37]. If this is equally true of rabbits, it may be that both agents use similar renal excretion and hepatic metabolism mechanisms that compete for the same receptor site, leading to the prolonged period of elimination of the drugs or their active metabolites.

The time to induction of 4.4 ± 0.8 min associated with AK in this study conforms with the time to the loss of righting reflex (5.1 min) in a similar acepromazine (2 mg.kg⁻¹) and ketamine (60 mg.kg⁻¹) anaesthesia reported by Amarpal et al. [3]. Also, this finding agrees with the previous similar study involving concurrent intramuscular administration of enrofloxacin (10 mg.kg⁻¹) and xylazine (5 mg.kg⁻¹)-ketamine (35 mg.kg⁻¹) anaesthesia which produced shorter time to induction (1.7 ± 0.1 min) but longer duration of recumbency (70.0 ± 6.0 min) than the control values of 3.4 ± 0.3 min and 62.0 ± 1.5 min respectively [2].

The mean HR in the AK group was mildly depressed for the first 50 min and proceeded to stabilize, except for a significant increase in HR at 10 min time intervals. This

observation was also reported in rabbits [3], cats [8], [39] and sheep [5]. Verstegen et al. [41], and Ingwersen et al. [26] did not find any significant changes in the mean HR after the administration of AK in cats, except an increase at the first 10 min time interval. Acepromazine-ketamine has also been reported to cause a significant increase in the HR in cats [40] and dogs [12]. These changes were due to ketamine sympathomimetic effect and possible vagolytic action [30], [31], [40], [42]. The apparent sparing effect of ketamine on the cardiovascular system, secondary to its sympathomimetic activity, was negated by acepromazine's sympatholytic properties as observed with AK group in this study. Although enrofloxacin alone is not known to have impact on the cardiovascular system [4], when combined with acepromazine-ketamine, it potentiates the cardiovascular depressive effect of acepromazine as shown in this study, especially towards the end of the trial. The decreased mean HR obtained with AKE throughout the trial period compared with the AK values still falls within the normal range accepted for awake rabbits [21], [23]. Thus, the drug induced decreased HR appears to be of no clinical significance in clinically healthy rabbits.

The increase in the mean respiratory frequency of the AKE group, from the 40 min time interval until the end of the study compared with the control values, is worthy of note. The mechanism for this could not be determined from this study. On its own, acepromazine at clinical doses has very little effect on respiration [19], [38]. The decrease in the respiratory frequency of the AK group in the first 40 min of our study can be attributed to the respiratory depressant properties of ketamine [19], [27], [29]. The decline in respiratory frequency was also reported in rabbits [25] and cats [8], [26] administered with acepromazine-ketamine. Enrofloxacin administered alone at therapeutic, or even excessive doses, does not exert significant effects on the vital parameters [4] and thus there is no expected side effect of acute nature in the clinics [6].

The significant difference in the mean RT with the AKE group than the AK group from 20th min time interval until the end of the study is interesting. All phenothiazine derivatives including acepromazine are known to cause a fall in body temperature, partly due to increased heat loss through dilated cutaneous vessels, and partly through resetting of thermoregulatory mechanisms [19], [28], [35], whereas ketamine is known to cause hyperthermia especially in cats [37]. In this study, the mean RT with AK group increased slightly for the first 40 min, but not significantly different from the baseline value and later stabilized until the end of the trial. This showed that the hypothermic effect of acepromazine counteracted the hyperthermic effect of ketamine. This becomes obvious when enrofloxacin potentiates the hypothermic effect of acepromazine, as observed in the AKE group until the end of the study.

In conclusion, the concurrent administration of enrofloxacin and acepromazine-ketamine anaesthesia in clinically healthy rabbits produced a shorter time to induction, but longer duration of recumbency and time to standing, than the corresponding control values. Significantly lower mean heart rate and rectal temperature, as well as, higher respi-

ratory frequency were also recorded compared to the control values. Therefore the use of this combination requires prolonged nursing care and caution in a cardiopulmonary compromised, hypovolaemic or critically ill patients, and also there is a need for body temperature monitoring and control in the AKE rabbits subjected to surgical exposure of the major body cavities, especially in a long duration procedure lasting more than one hour. However, it remains a good anaesthetic regimen in the healthy rabbit.

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EXAMPLES

Ahlborg, B., Ekelund, L. C., Nilsson, C. G., 1968: Effect of potassium-magnesium aspartate on the capacity of prolonged exercise in man. *Acta Physiol. Scand.*, 74, 238—245.

Black, H., Duganzich, D., 1995: A field evaluation of two vaccines against ovine pneumonic pasteurellosis. *New Zeal. Vet. J.*, 43, 60—63.

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FOLIA VETERINARIA, 57, 2013

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