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Citation for published version:

Aprich, V, Licka, T, Zipfl, N, Tichy, A & Gabriel, C 2017, 'Equine Hoof Canker: Cell Proliferation and Morphology' *Veterinary Pathology*, vol. 54, no. 4, pp. 661-668. DOI: 10.1177/0300985817695515

Digital Object Identifier (DOI):

[10.1177/0300985817695515](https://doi.org/10.1177/0300985817695515)

Link:

[Link to publication record in Edinburgh Research Explorer](#)

Document Version:

Peer reviewed version

Published In:

Veterinary Pathology

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Equine hoof canker: cell proliferation and morphology

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Abstract

Hoof canker is described as progressive pododermatitis of the equine hoof with absent epidermal cornification and extensive proliferation of the dermal papillary body; however, in-depth research on the type of proliferative activity has not yet been reported. The aim of the present study was to determine cell-specific proliferation patterns together with morphological analysis of hoof canker tissue. Tissues removed during surgery from 19 horses presented for treatment of canker were compared to similar postmortem tissues of healthy hooves of 10 horses. Morphological alterations visible in light microscopy were assessed semi-quantitatively and graded for severity. Proliferative activity was evaluated by means of anti-PCNA (proliferative cell nuclear antigen) and anti-Ki67 immunohistochemistry. Histologically, canker tissue showed five major morphological alterations, the presence of lacunae, vacuoles, giant cells, hemorrhage and inflammation, not seen in control tissue. Also, there was a notable koilocytotic appearance of keratinocytes in canker tissue. Immunohistochemistry revealed increased levels of PCNA protein expression in keratinocytes and fibroblasts of canker tissue compared to control tissue. In control tissue, keratinocytes showed higher levels of Ki67 compared to canker tissue, while the dermal fibroblasts of both groups showed similar levels of Ki67, indicating similar proliferative activity of less than 3% of total dermal fibroblasts. These results demonstrate that, in contrast to previous reports, there is no evidence for increased proliferative activity of the dermal papillary body associated with hoof canker. Increased levels of PCNA protein expression and morphological alterations indicate dysregulation of keratinocyte differentiation constitutes a key event in equine hoof canker development.

Key words: horse, hoof, canker, histology, immunohistochemistry, ki67, PCNA

For Peer Review

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3 Equine hoof canker (*Pododermatitis chronica verrucosa sive migrans*) is described
4
5 as destructive and progressive pododermatitis with alterations in the cornification
6
7 process of the keratinocytes (parakeratosis and hyperkeratosis).^{15, 24, 26} Extensive
8
9 proliferation of the dermal papillary body with subsequent absence of epidermal
10
11 cornification is reported to be a major feature of disease.^{23, 24} Hoof canker can affect
12
13 one or more hooves, and compromises the use and welfare of horses due to ensuing
14
15 instability of affected hoof capsules and lameness at late stages. Today, its etiology
16
17 is still unknown. Different theories for the development of hoof canker have been
18
19 reported. Inadequate husbandry, i.e. a lack of hoof hygiene and excessive moisture,
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21 as well as genetic predispositions, (auto)immune reactions, microorganisms (such as
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23 fungi, spirochetes and other anaerobic bacteria), as well as bovine papillomaviruses
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25 have been suggested to be involved in the development of canker.^{3, 9, 19, 20, 28}
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28 However, for any of these, a causal association with the development of disease
29
30 remains to be established. Morphological studies of hoof canker tissues are rare.^{24, 26}
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32
33 Ballooning of the keratinocytes, mainly in the Stratum germinativum of the frog, has
34
35 been reported together with the occurrence of perinuclear vacuoles in the cells of
36
37 deep layers of the Stratum spongiosum; the cells of the corium are covered by a thin
38
39 layer of defective epidermis, leading to an increased risk of infection and bleeding.²⁶
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41
42 To our knowledge, the proliferative character of canker has not yet been analyzed in
43
44 detail. Immunohistochemistry allows characterization of tissue proliferation, as has
45
46 been shown in a number of potentially metaplastic or neoplastic tissues.^{4, 13, 14, 27}
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49 During the proliferative phase of eukaryotic cells, different markers are used to
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51 determine the proliferative activity in healthy versus pathologically altered tissue.^{4, 13,}
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54 ²⁷ In the present study, antibodies to detect the Ki67 protein and the proliferating cell
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56 nuclear antigen (PCNA) protein were selected to analyze the proliferative processes
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3 in canker tissue. Ki67 is a sensitive, well-established predictive marker for
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5 recurrence in human breast cancer; it is widely used in tumor diagnostic and cell
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7 proliferation research.^{14, 27} PCNA is a marker for proliferation, and, additionally, it
8
9 indicates DNA damage and related repair mechanisms, but can be induced also by
10
11 growth factors independent of the cell cycle.¹⁴ Based on the hypothesis that equine
12
13 hoof canker is a proliferative disease, the aim of the present study was to investigate
14
15 the proliferative status and morphological alterations in canker tissue compared to
16
17 healthy hoof tissue.
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20 21 22 **Materials and Methods**

23 24 **Sample collection**

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26
27 Tissues of 19 horses presented for treatment of hoof canker to the Equine University
28
29 Clinic of the University of Veterinary Medicine, Vienna, Austria (Supplemental Table
30
31 1) were assessed. Hoof canker was diagnosed based on its typical clinical
32
33 appearance.^{1, 23, 24} Tissue samples were taken from superficial regions of canker
34
35 affected areas with a hoof knife on the first day of treatment and from deep regions
36
37 including the dermo-epithelial junction during surgery under regional anesthesia on
38
39 the following day. Tissue sampling and evaluation, as well as anonymized
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41 publication of the data obtained, were carried out with the written consent of the
42
43 horse owners, given on admission of the horse to the clinic. Sample collection was
44
45 carried out during therapeutic trimming and surgery of affected hooves; all tissues
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47 used were removed based on clinical necessity.
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52 Samples of 11 horses euthanized for reasons other than hoof or skin disease were
53
54 used as control tissue (Supplemental Table 1). Prior to inclusion as controls, all
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56 hooves and areas of skin were examined macroscopically and histologically for
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3 absence of pathological alterations. Therefore, hoof tissues of corresponding regions
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5 of 10/11 horses were sampled post mortem, and skin of 1/11 horse was used for the
6
7 establishment of immunohistochemical (IHC) protocols.
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10 11 Histology and Immunohistochemistry

12
13 Tissues were cut into 1 x 2 cm pieces for histological embedding. Tissues for
14
15 histological and IHC analysis were immersion-fixed in 4% buffered formaldehyde for
16
17 a minimum of 72 hours and embedded in Paraplast® (Vogel, Giessen, Germany).
18
19 Histological sections of 3µm thickness were cut and stained with haematoxylin and
20
21 eosin (H&E) according to Romeis for morphological analysis.¹⁶ For IHC serial
22
23 sections (3 µm) of skin and hoof, tissues were mounted on 3-
24
25 aminopropyltriethoxysilane/glutaraldehyde - coated slides. Endogenous peroxidase
26
27 activity was blocked by incubation in 0.6% H₂O₂ in methanol for 15 min at room
28
29 temperature. A protein block (1.5% normal goat serum) was used to minimize
30
31 unspecific binding of the primary antibody. The unlabeled monoclonal primary
32
33 antibodies (anti-PCNA and anti-Ki67; for sources, pre-treatments and dilutions - see
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35 Supplemental Table 2) were detected with the ImmunoVision secondary system
36
37 (ImmunoVision Technologies, Brisbane, CA, USA) using DAB (3,
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39 3'-diaminobenzidine-tetrahydrochloride substrate in Tris buffer pH 7.4 and 0.03%
40
41 H₂O₂) as chromogen. Finally, slides were washed with distilled water, counterstained
42
43 with haemalum, dehydrated and mounted by use of xylene-soluble medium (DPX,
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45 Fluka, Buchs, Switzerland). Negative controls were obtained by substitution of the
46
47 primary antibodies by PBS. Evaluation of the sections was performed using light
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49 microscopy (Polyvar, Reichert-Jung, Vienna, Austria) and a digital camera (Nikon,
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51 Austria). Sections of healthy equine skin were used as positive controls.
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Morphological analysis

Three H&E-stained histological sections per tissue sample and two to five tissue samples per horse were evaluated. Five frequently occurring alterations of canker tissue were analyzed and evaluated in a semi-quantitative setup (Table 1). The grading was performed in a blinded way by one person and combined frequency and intensity of the respective criteria. The alterations were graded for intensity as mild (one point), moderate (two points) or severe (three points) and for frequency concerning the occurrence of the respective criterion in less than 20% of the examined area (one point), between 20 and 50% (two points) and more than 50% (three points). The sum of the grades of all investigated sections per horse was calculated and the mean value for each criterion was determined per horse.

Lacunae were defined as fluid filled cavities with eosinophilic content, vacuoles as formations surrounding the nucleus of keratinocytes in different manifestations, giant cells presented as enlarged keratinocytes with morphologically intact cytoplasm and nucleus, hemorrhage was described as erythrocyte accumulation and tissue based neutrophils, lymphocytes and monocytes indicated inflammation.

Scoring of Proliferative Activity

Proliferative activity was assessed using two antibodies (anti-Ki67, anti-PCNA). The nuclei positive for the respective protein were counted in three randomly selected fields featuring a minimum of 100 cells per field on two different slides for each tissue sample. In three different regions (Stratum basale, Stratum germinativum, and the dermal portion at the dermo-epithelial transition zone in the papillae) proliferative activity was determined as the number of antibody positive cells as a percentage of

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3 total cell number counted per field for each antibody (scoring index). In the present
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5 study, the Stratum germinativum was defined as the entire proliferative zone of the
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7 hoof epidermis, combining the single-layered Stratum basale and the proliferative
8
9 cells of the Stratum spinosum.
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11 12 13 14 Statistical analysis

15
16 A linear mixed model using SPSS statistics 20.0 software (IBM; Vienna, Austria) was
17
18 applied to identify significant differences of proliferative activity (determined via
19
20 PCNA and Ki67 protein expression, respectively) between healthy and hoof canker
21
22 affected cell types of the respective regions. Furthermore, a linear mixed model
23
24 followed by Sidak correction was adapted to compare the values of proliferative
25
26 activity between the different regions for each group (hoof canker tissue and control
27
28 tissue). For all statistical tests a p-value ≤ 0.05 was considered significant.
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32 33 34 **Results**

35 36 Clinical findings

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38 Ten geldings and nine mares (Supplemental Table 1) presenting for hoof canker at
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40 the Equine University Clinic of the University of Veterinary Medicine, Vienna, Austria,
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42 from 2011 to 2014 were included in this study. The average age was 15 years (7 –
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44 24 years). In 10/19 horses only one hoof showed canker, while in 6/19 horses two
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46 and in 3/19 horses three hooves were affected. Hoof canker was diagnosed in hind
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48 limbs in 9/31 hooves, whereas front limbs were affected in 22/31 hooves
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50 (Supplemental Table 1).
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55 56 Morphological analysis

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3 Results of the morphological evaluation of histological sections obtained from hoof
4
5 canker tissue are listed in table 1. The presence of lacunae was mainly observed at
6
7 the end of papillae (Figures 1-3). In mildly affected tissue, vacuoles were detected as
8
9 slim white halos surrounding intact nuclei as revealed by H&E staining. In severely
10
11 affected tissue, nuclei were pyknotic and cells were enlarged and filled with
12
13 perinuclear vacuoles. These cells were referred to as ballooning keratinocytes or
14
15 koilocytotic keratinocytes (Figures 1, 3, 5 and 7). Giant cells (Figure 6) were
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17 analyzed by frequency only. Hemorrhage in the epidermal tissue was identified on
18
19 basis of erythrocyte accumulation. In all canker tissues signs of inflammation
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21 including tissue based neutrophils, lymphocytes and monocytes were present in
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23 varying severity (Figures 4 and 6).
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29 Proliferative Activity

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31 Immunohistochemical analyses of PCNA and Ki67 protein expression showed layer
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33 specific differences (Figures 8-11). In control tissue, epidermal PCNA protein
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35 expression was of lower intensity and frequency in the Stratum basale and the
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37 Stratum spinosum compared to corresponding canker tissue (Figures 8 and 9). In
38
39 the hoof canker tissue, a high number of PCNA-expressing keratinocyte nuclei were
40
41 located in the Stratum spinosum. In the affected dermis fibroblasts frequently
42
43 showed PCNA expression in the nuclei; this was a less common observation in
44
45 control tissue. The Ki67 protein expression patterns were distinct from PCNA
46
47 expression patterns in both sets of tissues in regard to reactivity and localization
48
49 (Figures 10 and 11). Keratinocyte nuclei expressing Ki67 were predominantly found
50
51 in the Stratum basale and the deep layers of the Stratum spinosum. Ki67-positive
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53 fibroblasts were rare in both sets of tissues. Negative controls showed no staining in
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3 all IHC reactions (Figures 12 and 13), but melanin granules were occasionally
4
5 observed in the Stratum basale and the Stratum spinosum.
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7 Results of quantitative analysis are summarized in supplemental table 3 and
8
9 presented in figures 14 and 15. The number of PCNA-positive keratinocytes in the
10
11 Stratum basale was significantly higher in canker tissue than in control tissue. A
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13 similar scenario was observed for the entire Stratum germinativum. In canker tissue,
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15 the numbers of PCNA-positive keratinocytes in the Stratum basale was significantly
16
17 lower compared to the Stratum germinativum. This significant difference was not
18
19 evident in the control tissue. The number of PCNA-positive dermal fibroblasts was
20
21 significantly smaller than the number of PCNA-positive epidermal cells in canker
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23 tissue, there was no significant difference in the control tissue.
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27 The Ki67 protein staining revealed only a small population of positive cells in canker
28
29 and control tissue. In control tissue, proliferative activity of the keratinocytes, as
30
31 determined by Ki67 staining, was higher than in canker tissue in the Stratum basale,
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33 as well as in the entire Stratum germinativum. In both sets of tissues, the scoring
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35 index for Ki67 in dermal fibroblasts was similar (< 3%).
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40 **Discussion**

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42 Despite the reported higher incidence of hoof canker in hind feet, more than two
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44 thirds of the affected feet were front feet in this study.^{20, 25} The main limitation of this
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46 study was that canker was diagnosed based on its macroscopic appearance in the
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48 absence of any other gold standard of diagnosis; main outcomes were microscopic
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50 features, therefore, introducing a potential risk of circular argumentation.
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54 In the present study, the monoclonal anti-Ki67 antibody detected fewer positive
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56 keratinocytes in canker tissue compared to control tissue. The protein Ki67 is a
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3 highly sensitive marker of proliferating cells in the G1, S and G2 phase as well as
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5 during mitosis, but it cannot be detected in cells in G0 phase.²⁷ In contrast, PCNA
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7 IHC staining resulted in significantly more positive keratinocytes in canker tissue
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9 compared to control tissue, illustrating either a higher proliferative activity or DNA
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11 damage-related repair mechanisms in canker tissue. The protein PCNA is more
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13 stably expressed and can also be detected at the end of the cell cycle; in its function
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15 as a nuclear ring clamp protein, PCNA is also involved in DNA repair mechanisms
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17 and can therefore be induced also by growth factors independently from the cell
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19 cycle.^{14, 33} The detection of elevated numbers of PCNA-positive cells and reduced
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21 numbers of Ki67-positive cells in the same samples, as found in the present study,
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23 has already been described in a variety of tumors and normal tissues.^{2, 6, 10, 17}
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27 Possible scenarios leading to such results include the detection of PCNA protein at
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29 the end of cell cycle, as well as in non-proliferating cells during DNA repair or growth
30
31 factor activation. Therefore, it is reasonable to find more PCNA positive cells than
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33 Ki67 positive cells. This also explains, why Ki67 protein is the more sensitive and
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35 specific marker for proliferative events in tumor diagnosis.^{6, 10, 17} Based on this
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37 knowledge, the Ki67 protein expression data in the present study served to
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39 conclude, that no increase of proliferative activity is observed for the epidermis and
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41 dermis of canker tissue.
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45 The protruding tissue, which is so typical for hoof canker, lacked keratinization and
46
47 related desquamation, whilst hypertrophic events abounded, as revealed by lacunae
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49 formation, ballooning keratinocytes and giant cell formation, demonstrated in the
50
51 present study. These hypertrophic events in combination with loss of tissue integrity
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53 might explain the typical filamentous or cauliflower-like appearance of hoof canker,
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55 as the dermal parts start to prolapse due to lack of pressure of the horn capsule; this
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3 is an effect often seen after the creation of hoof horn defects, if too little pressure is
4 exerted onto the granulation tissue.³⁶ Lack of keratinization becomes apparent
5 through epidermal cells, which remain spherical, thus creating a larger tissue volume
6 without hyper-proliferating. In addition, the production of abnormal quantities or types
7 of cell proteins possibly contributes to the clinical picture of extensive growth and
8 white cheesy exudate.^{5, 21}

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16 In contrast to other descriptions of canker, the present study provides no evidence of
17 increased proliferative activity.^{19, 30} Yet, hypertrophy and DNA damage with
18 subsequent disturbance of keratinization of epidermal cells were frequently
19 encountered.

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25 Not surprisingly, hemorrhage was observed in 18 of 19 surgical hoof canker
26 samples. The hemorrhage was potentially enhanced by detected inflammatory
27 processes also present in all canker tissues. Inflamed regions were commonly found
28 in and adjacent to the dermal papillae, potentially induced by reduced tissue
29 cohesion between maldifferentiated, not cornified epidermis and corium. A high
30 number of free connective tissue cells were observed in the dermal portion of the
31 dermo-epithelial zone and in the dermal papillae as additional sign of inflammation.
32 Some samples contained bacteria, as the barrier function of the epidermis is
33 impaired with the cessation of keratinocyte differentiation. This is similar to previous
34 reports, that bacterial infection is an inconsistent finding in hoof canker.³¹

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47 In most hoof canker samples, formation of large epidermal lacunae was observed;
48 these lacunae were filled with eosinophilic fluid and might result from accumulation
49 or increased production of this fluid (Figure 2). Lacunae formation is suggested to be
50 related to instability of the epidermal tissue.^{18, 32} The instability of the hoof capsule is
51 a common cause for canker-associated lameness in late stages of the disease.³⁰

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3 Hoof canker-associated keratinocytes are commonly described as “ballooning
4
5 keratinocytes” because of higher cell volumes, pyknotic nuclei and perinuclear
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7 vacuoles, which was also found in the present study.^{19, 24} In human keratinocytes,
8
9 human papillomavirus type 16 (HPV-16)-induced changes include comparable
10
11 alterations of the keratinocytes (and human cervical cells in vitro) described as
12
13 koilocytotic cells. These cells feature perinuclear vacuoles appearing as white halos
14
15 and mostly acentric, hyperchromatic nuclei.^{11, 12} The same cellular appearance is
16
17 also reported for other virus-infected cells, including *Molluscum Contagiosum*, a
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19 poxvirus infection in humans, leading to cutaneous lesions with acanthosis,
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21 hyperkeratosis and nuclear atypia including koilocytotic keratinocytes.²⁹ Therefore,
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23 koilocytosis observed in keratinocytes of equine hoof canker tissue may refer to a
24
25 potential viral involvement in etiology. The success of local cisplatin chemotherapy
26
27 for equine hoof canker is comparable to the success of intralesional cisplatin
28
29 chemotherapy for papillomaviral-induced equine sarcoids, which is a well-
30
31 documented therapy for this disease.^{1, 7, 34, 35} In addition, the increased expression of
32
33 PCNA protein in hoof canker-associated keratinocytes is another parallel with
34
35 symptoms of viral infection, as, in human keratinocytes in vitro, PCNA expression is
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37 down-regulated by growth inhibition via TGF β or calcium induced differentiation,
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39 while in HPV transformed keratinocytes this effect was reduced and PCNA
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41 expression was still high.⁸ These observations support the concept of
42
43 papillomaviruses being involved in the development of equine hoof canker, although
44
45 the causal association with onset and progression still remains to be established.³
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47 Still, neither morphological analyses, nor detected proliferative activity of hoof canker
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49 tissue revealed evidence of neoplastic transformation of cells of the papillary body. In
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51 differentiated keratinocytes reduced PCNA levels were found, which might also
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3 explain the significantly higher PCNA levels in the poorly differentiated keratinocytes
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5 in hoof canker.^{8, 22} Together with the acanthosis seen in most samples, this is
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7 additional evidence that hoof canker is related to disturbances in differentiation of the
8
9 keratinocytes rather than a proliferative dysfunction.
10

11 The results of the present study suggest that hoof canker is not a hyperproliferative
12
13 process of the dermal papillary body shown by immunohistochemical anti-Ki67
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15 detection, but increased levels of PCNA protein expression and observed
16
17 morphological alterations indicate that dysregulation of keratinocyte differentiation
18
19 constitutes a key event in equine hoof canker development.
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25 **Acknowledgements:** The authors give thanks to Claudia Höchsmann and Stefanie
26
27 Burger for the establishment of the immunohistochemical staining protocols and to
28
29 Birgit Machac and Anne Flemming for assisting in production of the histological
30
31 sections.
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37 **Figure legends**

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39 Figures 1-7. Hoof canker, equine, hematoxylin and eosin stain. Figure 1: Enlarged,
40 fluid filled lacunae (L) are surrounded by enlarged keratinocytes with an increased
41 amount of lightly eosinophilic to clear cytoplasm surrounding the nucleus (ballooning
42 degeneration of the keratinocytes). Figure 2: Lacunae (L) are encircled by a thin
43 membrane-like structure (arrows). Figure 3: Normal keratinocyte nuclei are
44 surrounded by a white halo and the cytoplasm contains variably sized eosinophilic
45 droplets. Beside intracellular localization, eosinophilic droplets can also be observed
46 within the lacunae. Figure 4: High numbers of inflammatory cells (mostly
47 eosinophilic and neutrophil granulocytes) were observed in the corium near the intact
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3 Stratum germinativum (SG). Figure 5: Regions of moderate ballooning degeneration
4 of keratinocytes (top and middle right hand side) are present near regions of normal
5 Stratum germinativum (SG) and regions around the dermal papillae (DP). Figure 6:
6 Scattered cells within the normal stratum germinativum were karyomegalic (giant
7 cells) and had a perinuclear clear halo (arrows). Figure 7: Area of clear demarcation
8 (arrows) between the normal Stratum germinativum (SG) and the keratinocytes with
9 ballooning degeneration.
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21 Figures 8-13. Immunohistochemical analysis of PCNA and Ki67 protein expression
22 in canker tissue and control hoof tissue. Figure 8: Canker tissue, PCNA protein was
23 mainly observed in epithelial cells of the Stratum basale (arrows), the whole of the
24 Stratum germinativum (SG) and in fibroblasts within the dermal papillae (DP). Figure
25 9: Control hoof tissue, strong anti-PCNA staining was observed in keratinocytes of
26 the Stratum basale (arrows) and in keratinocytes of the Stratum germinativum. In the
27 dermal papillae fibroblasts were negative for PCNA. Figure 10: Canker tissue, Ki67
28 protein expression was restricted to the Stratum basale (arrows) and the deep layers
29 of the Stratum spinosum within the Stratum germinativum (SG). Scattered fibroblasts
30 were positive for anti-Ki67 staining in the dermal papillae (DP). Figure 11: Control
31 tissue, keratinocytes were positive for the Ki67 protein in the Stratum basale (arrow)
32 and the deep layers of the Stratum spinosum within the Stratum germinativum (SG),
33 whereas a smaller number of dermal fibroblasts were positive in the dermal papillae
34 (DP). Figures 12 and 13: Hoof canker, negative controls for the anti-PCNA (Figure
35 12) and anti-Ki67 (Figure 13) immunohistochemical stain. Melanin granules (arrows)
36 were occasionally observed in the Stratum germinativum.
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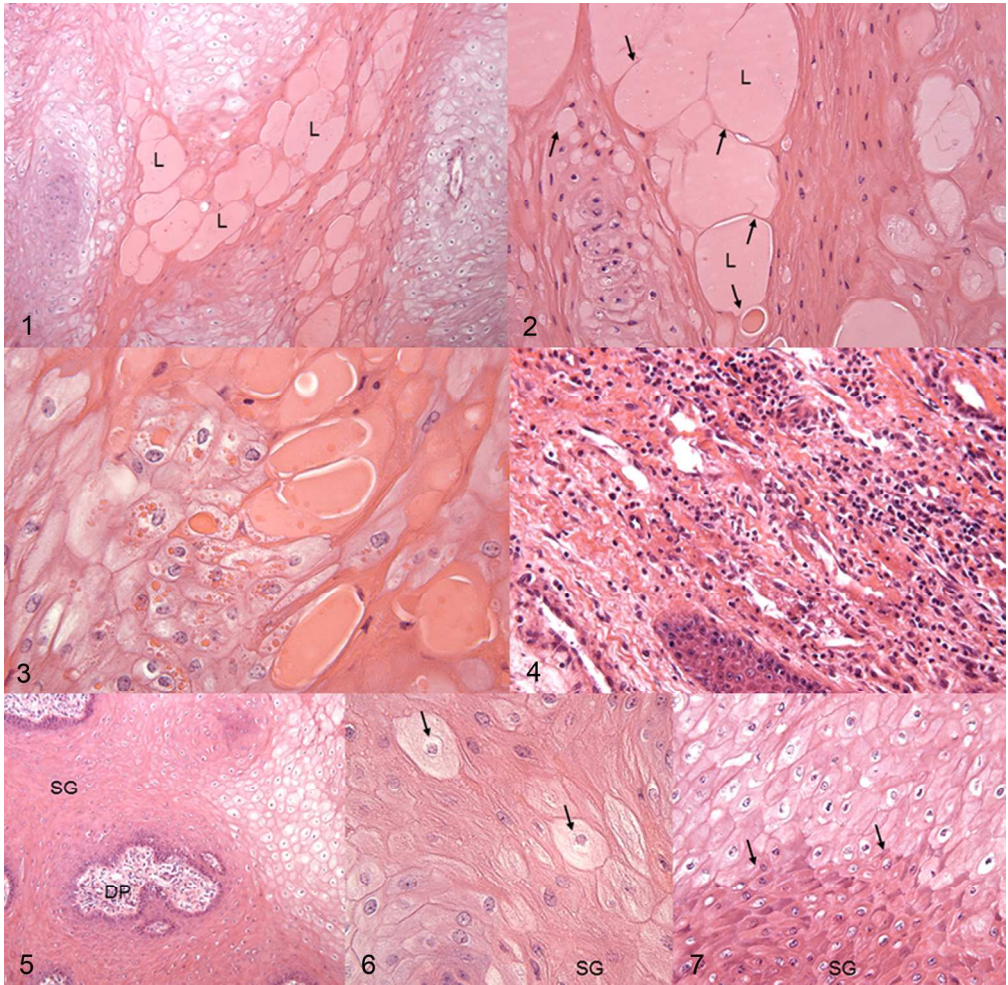
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3 Figures 14 and 15. Evaluation of the percentage of anti-PCNA (Figure 14) and anti-
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5 Ki67 (Figure 15) positive nuclei in the keratinocytes of the Stratum basale and the
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7 total Stratum germinativum, and the fibroblasts of the dermal connective tissue in
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9 cancer samples (n = 19) and control samples (n = 10). Identical characters (a - e)
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11 indicate significant differences ($p \leq 0.05$) between the groups and regions.
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For Peer Review

Table 1: Summary of the semi-quantitative evaluation of morphological alterations performed on histological sections of 19 hoof canker affected horses; the grading combined frequency and intensity of the respective criterion.

Horse	Lacunae	Vacuoles	Giant cells	Hemorrhage	Inflammation
1	-	+	+	++	++
2	+	+	+++	++	++
3	+	++	+++	++	++
4	-	-	-	-	+
5	++	++	+++	++	++
6	+	++	++	+	++
7	++	+++	+++	+	++
8	+	++	++	++	++
9	++	++	+	+	+
10	++	+++	++	+	++
11	+	++	+++	++	++
12	+	++	++	++	+
13	++	+++	+++	+	++
14	+++	++	++	+	++
15	++	++	++	+	++
16	+	++	+++	++	++
17	+	+++	+++	++	+++
18	-	++	++	+	++
19	++	+++	++	+	++

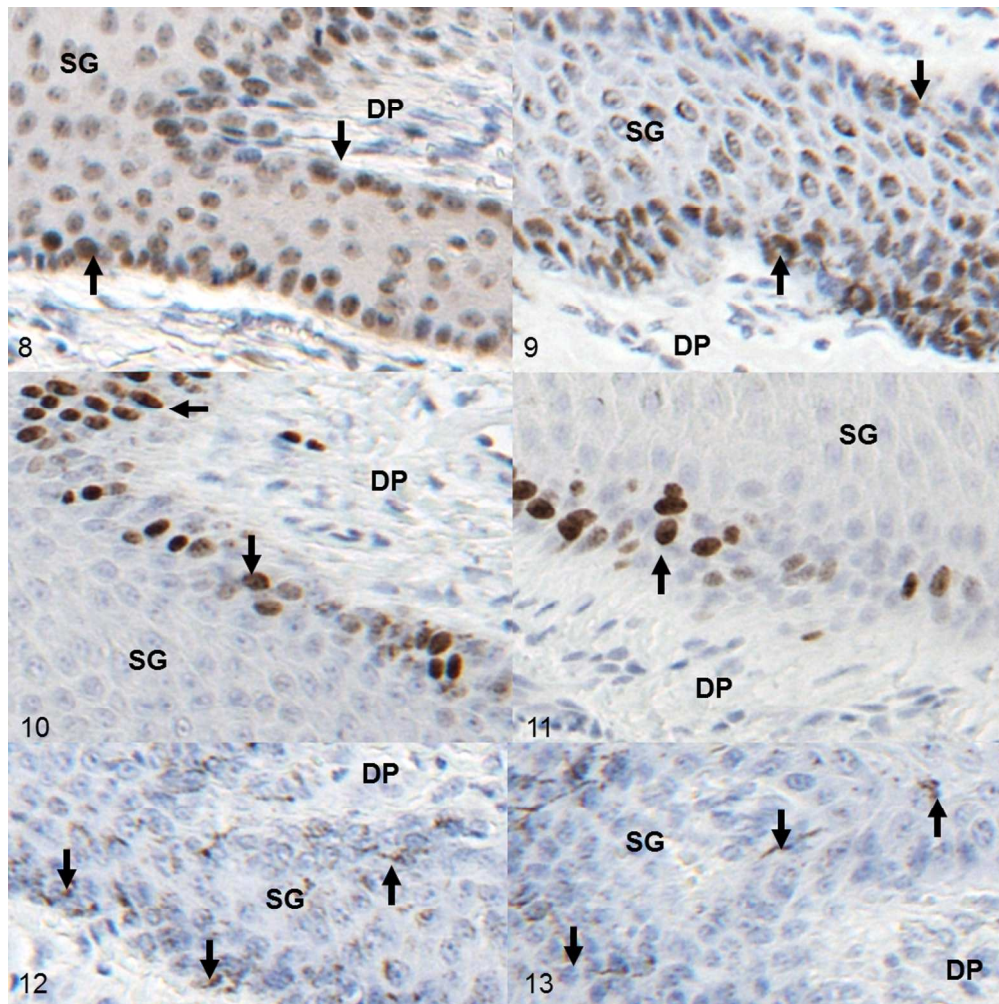
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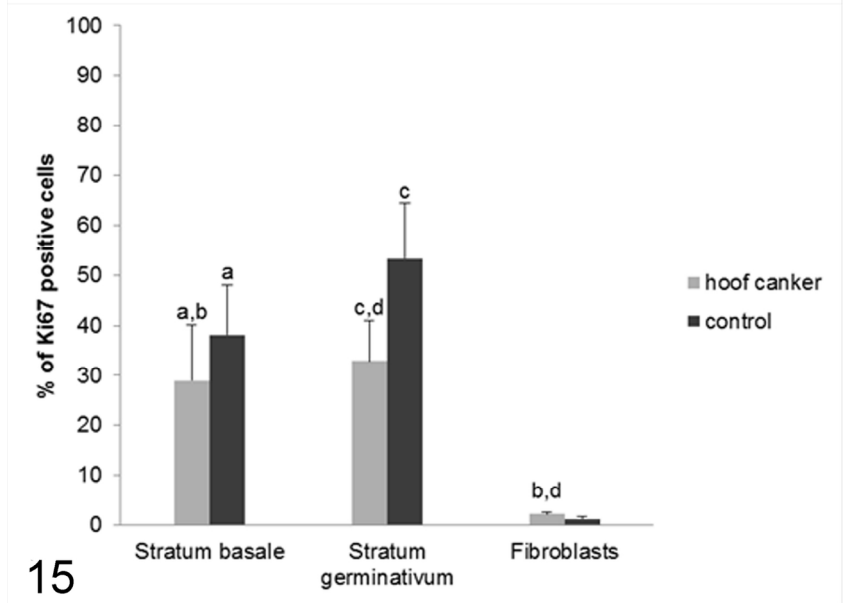
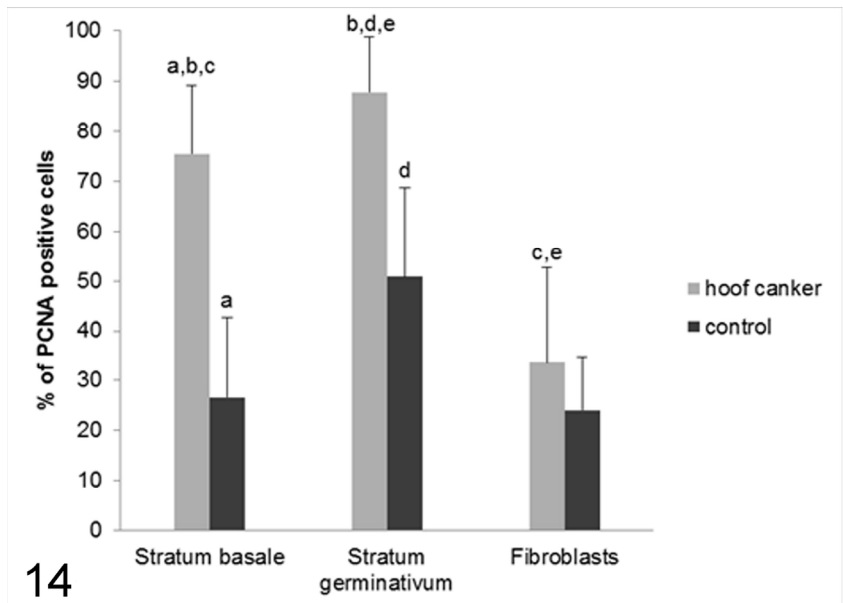
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Supplemental table 1: Clinical data of hoof canker affected horses (1-19) and control horses (20-30) presented in this study. G gelding; M mare; S stallion; R right; L left; H hind hoof; F front hoof

horse	sex	age (years)	breed	hoof/hooves
1	G	9	Warmblood	RH
2	G	13	Holsteiner	LF
3	G	15	Pinto	LF
4	G	13	Holsteiner	LH, RH
5	M	15	Noriker	LF, RF, RH
6	M	18	Arabian	LH
7	M	15	Warmblood	LF
8	M	12	Trotter	LF, RF
9	G	8	Kladruber	LF, LH, RH
10	G	16	Warmblood	LF
11	M	16	Haflinger	LF, RF
12	M	24	Trotter	LF, RF
13	G	18	Warmblood	LF, RF, LH
14	M	11	Noriker	LF
15	M	16	Trotter	RF
16	G	15	Hanoverian	LF, RF
17	G	7	Warmblood	RH
18	G	20	Trotter	LF, RF
19	M	18	Trotter	RF
20	M	14	Warmblood	LF
21	M	16	Shagya Arabian	RF, RH
22	G	7	Trotter	LH
23	S	8	Noriker	RH
24	M	14	Haflinger	RH
25	S	2	Lusitano	RH
26	G	6	Lusitano	LF, RF
27	G	17	Warmblood	LH, RH
28	M	4	Quater Horse	RF
29	M	6	Warmblood	LF
30	M	17	Tinker	skin

Supplemental Table 2: Sources, pre-treatments and dilutions of the antibodies used in this study.

<u>Antibody</u>	<u>Source</u>	<u>Clone</u>	<u>Pretreatment</u>	<u>Dilution</u>
anti-Ki67	Thermo Fisher Scientific, Fremont, CA, USA	SP6 rabbit	citrate buffer, pH 6.0, 2 hours at 85°C	1 : 400
anti-PCNA	NeoMarkers, Fremont, CA, USA	PC10 mouse	citrate buffer, pH 6.0, 2 hours at 65°C	1 : 1000

Supplemental table 3: Relative amount of positive cells stained for PCNA and Ki67 protein by means of immunohistochemistry in three different regions of hoof canker affected horses (n = 19) and the control group (n = 10). The values are mean values \pm SDs.

	Scoring index PCNA		Scoring index Ki67	
	Hoof canker	Control	Hoof canker	Control
Stratum basale	75.55 \pm 27.68	26.62 \pm 16.05	28.93 \pm 22.54	38.13 \pm 9.88
Stratum germinativum	87.90 \pm 21.96	50.88 \pm 17.80	32.96 \pm 16.08	53.32 \pm 11.21
Fibroblasts (Corium)	37.75 \pm 29.86	24.07 \pm 10.73	2.22 \pm 0.71	1.20 \pm 0.45