Characterization of equine hyalocytes: their immunohistochemical properties, morphologies and distribution

Yuto SANO¹, Kazuya MATSUDA¹, Minoru OKAMOTO¹, Kazushige TAKEHANA², Kazuko HIRAYAMA¹ and Hiroyuki TANIYAMA¹

¹Department of Veterinary Pathology, School of Veterinary Medicine, Rakuno Gakuen University, Ebetsu, Hokkaido 069–8501, Japan
²Department of Veterinary Microanatomy, School of Veterinary Medicine, Rakuno Gakuen University, Ebetsu, Hokkaido 069–8501, Japan

(Received 1 September 2015/Accepted 2 February 2016/Published online in J-STAGE 18 February 2016)

KEY WORDS: ciliary body, horse, hyalocytes, vitreous cortex

In horse, the characterizations of hyalocytes under the steady state are still unclear. Therefore, we investigated characterizations of hyalocytes in normal equine eyes by their immunohistochemical phenotype, histomorphology and distribution. Thirty-one eyes from 18 horses, divided into 4 groups (G) by age, were used: early (G1) and late gestation (G2) fetuses, 1- to 3-year-old (G3) and 8- to 24-year-old (G4) horses. Equine hyalocytes were histologically classified into 4 types, and they immunohistochemically expressed MHC II and CD163. Hyalocytes were detected on and/or around ciliary processes and pars plana in G2, G3 and G4, but were not located on retina and optic papilla. A significant increase in distribution was found between G2 and both G3 and G4, and the largest distribution was found at ciliary processes in these groups. Equine hyalocytes were characterized as residential ocular macrophage and MHC II antigen-bearing cell, accompanied by a pleomorphic appearance and located in the contiguous ciliary body. Our data provided characterizations of hyalocytes in normal equine eyes and may well contribute to improving the understanding of pathogenesis of equine ocular disease.

MATERIALS AND METHODS

Animals: Thirty-one normal eyes from 18 horses were studied. The horses used in this study were submitted for necropsy to the Laboratory of Veterinary Pathology of Rakuno Gakuen University. The breeding distribution was thoroughbred. The age of the horses ranged from about 90 days of fetus gestation to 24 years of age. There were a variety of clinical and pathological diagnoses, but ocular abnormalities were not observed in any horses. Horses used in this study were divided into 4 groups by ages: 6 eyes from early gestation fetuses (Group 1; G1), 6 eyes from late gestation (Group 2; G2) fetuses, 8 eyes from 1- to 3-year-old horses (Group 3; G3) and 11 eyes from 8- to 24-year-old (Group 4; G4) adult horses. The characterizations of horses used in this study and their divided groups are summarized in Table 1.

Tissue preparation: The fixative solution used in the present study was 10% neutral buffered formalin. For adequate fixation, appendages of the eyes were trimmed, and an incision of about 1 cm was made dorsal and sagittal ora serrata. The formalin was injected from the incision, and then, the eyes were immersed for 24 to 48 hr in formalin. Fixed eyes were divided into anterior and posterior ocular segments. The divided anterior ocular segment was cut into pieces of two or three 0.5–10 mm strips of tissue containing cornea, iris and ciliary body. The divided posterior ocular segment was also cut into two or three 0.5–10 mm strips of tissue containing retina and optic papilla. The tissues were embedded in paraffin wax following the immersion of graded ethyl alcohols and xylene. The embedded ocular tissues were sec-
Table 1. Characterizations of horses used in the present study

<table>
<thead>
<tr>
<th>Group</th>
<th>Eye</th>
<th>Sex</th>
<th>Age</th>
<th>Diagnosis</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>R/L</td>
<td>M</td>
<td>90 d</td>
<td>—</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>F</td>
<td>90 d</td>
<td>—</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>M</td>
<td>90 d</td>
<td>—</td>
</tr>
<tr>
<td>2</td>
<td>R/L</td>
<td>M</td>
<td>310 d</td>
<td>Difficult delivery and stillborn</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>M</td>
<td>318 d</td>
<td>Difficult delivery and stillborn</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>M</td>
<td>320 d</td>
<td>Difficult delivery and stillborn</td>
</tr>
<tr>
<td>3</td>
<td>R/L</td>
<td>M</td>
<td>1 y</td>
<td>Osteochondritis disseccans</td>
</tr>
<tr>
<td></td>
<td>L</td>
<td>M</td>
<td>1 y</td>
<td>Wobbler syndrome</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>F</td>
<td>2 y</td>
<td>Wobbler syndrome</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>M</td>
<td>2 y</td>
<td>Wobbler syndrome</td>
</tr>
<tr>
<td>4</td>
<td>R/L</td>
<td>F</td>
<td>8 y</td>
<td>Colonic torsion</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>F</td>
<td>8 y</td>
<td>Rupture of adductor muscle</td>
</tr>
<tr>
<td></td>
<td>R</td>
<td>F</td>
<td>9 y</td>
<td>Chronic laminitis</td>
</tr>
<tr>
<td></td>
<td>L</td>
<td>F</td>
<td>9 y</td>
<td>Spinal cord injury</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>F</td>
<td>15 y</td>
<td>Chronic laminitis</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>F</td>
<td>24 y</td>
<td>Squamous cell carcinoma</td>
</tr>
<tr>
<td></td>
<td>R/L</td>
<td>F</td>
<td>24 y</td>
<td>Mammary gland adenocarcinoma</td>
</tr>
</tbody>
</table>

R: right, L: left, M: male, F: female, d: day, y: year-old.

Immunohistochemical examinations: Indirect immunofluorescence studies were performed, and the primary antibodies against MHC class II (MHC II), CD163, CD3 and CD20 were used to recognize antigen-presenting cells, monocyte/macrophages and lymphocytes (T and B cells), respectively. Glial fibrillary acidic protein (GFAP) and pan-cytokeratin were also used to differentiate glial cells and epithelial cells, and the S100 protein was used as a functional marker for Ca²⁺ binding protein. The details of the primary antibodies and pretreatments for antigen retrieval are listed in Table 2. After pretreatment, the sections were incubated with 10% normal goat serum (Sigma Chemical Co., St. Louis, MO, U.S.A.) at 37°C for 30 min as a blocking step. Subsequently, the sections with primary antibodies were incubated at 4°C overnight in moist chambers. Following the primary incubation, the sections were reacted with Alexa Fluor 488-labeled goat anti-mouse IgG (Molecular Probes, Eugene, OR, U.S.A., diluted 1:200) or Alexa Fluor 546-labeled goat anti-rabbit IgG (Molecular Probes, diluted 1:200) at room temperature for 30 min in shaking moist chambers. The sections were covered with water-soluble mounting medium and then were examined using a confocal microscope (C2; Nikon Instech Co., Ltd., Tokyo, Japan). With each immunofluorescence run, equine lymph node was used as a positive control for the detection of MHC II, CD163, CD3 and CD20. For the detection of GFAP and S100, and pan-cytokeratin, equine retina and epidermis were also used as positive controls, respectively.

Morphometry and statistical analyses: Hyalocytes from each separate ocular segment were manually counted: ciliary processes, pars plana, retina and optic papilla. Counting of the cells was performed in five different fields under a HE stained microscope image using 400× high-power magnifications. The length of the vitreous cortex surface in each image was measured by using public software (Image J). The counted number of hyalocytes was presented as the average per 500 μm. Statistical analyses were performed using specific software (Excel and Ekuseru-Toukei 2012; Social Survey Research Information Co., Ltd., Tokyo, Japan). To determine the distribution of hyalocytes among different ages, Kruskal-Wallis test and Scheffe’s test were used for a priori and post hoc comparison, respectively. To determine the site of the greatest distribution, the number of sites in each group was compared using Mann-Whitney U test. For each analysis, P<0.01 or P<0.05 was considered to be significant.

RESULTS

Histological findings: All HE-stained equine ocular sections used in this study were assessed by microscopic examination, and no abnormalities were recognized. In G3 and 4, hyalocytes located in the vitreous cortex were significantly detected in ciliary processes (Fig. 1) and pars plana. The cells were pleomorphic and were classified according to their 4 different morphological features: round or oval nuclei and cytoplasm (type-1, Fig. 2a), round or oval nuclei and cytoplasm with short or long processes (type-2, Fig. 2b), flattened or elliptical nuclei and flattened or elongated-shaped cytoplasm (type-3, Fig. 2c), and flattened or elliptical nuclei and flattened or elongated-shaped cytoplasm with short or long processes (type-4, Fig. 2d). The cells of type-2 and type-4 were more detectable than those of type-1 and type-3. The free cells of type-1 and type-2 were detected around ciliary processes. In G2, fewer hyalocytes were randomly located in the vitreous cortex of ciliary processes and pars plana. The 4 different morphological features were also detected, but the cells of type-1 and type-3 were more detectable than type-2 and type-4. The free cells of type-1 and type-2 were also detected around ciliary processes. In G1, the cells were not detected on any part of vitreous cortex. Hyalocytes were not found in the vitreous cortex of retina and optic papilla in any groups.

Immunohistochemical findings: All 4 types of hyalocytes in G2, G3 and G4 expressed MHC II (Fig. 3a and 3b) and CD163 (Fig. 4a and 4b). The free cells around ciliary processes also expressed each antibody. These cells did not express CD3, CD20, GFAP, S100 and pan-cytokeratin. In G1, the expressed cell of all antibodies was not detected in the vitreous cortex.
Hyalocytes are not differentiated to lymphocytes, glial cells CD20, GFAP and pan-cytokeratin suggested that equine ing cells [26]. The negative immnoreactive results for CD3, II antigen-bearing cells are considered to be antigen-present-
function in antigen presentation, as evidenced by MHC II expression usually can be identified in
in liver [17, 18, 31]. Moreover, equine hyalocytes have a
in spleen, alveolar macrophages in lungs and Kupffer cells
macrophages in normal tissues, such as interfollicular and
specifically identified in subpopulations of resident tissue
receptor that is selectively expressed in the cells belonging
to the monocyte/macrophage lineage [9]. The expression is
specifically identified in subpopulations of resident tissue
macrophages in normal tissues, such as interfollicular and
sinus macrophages in lymph nodes, red pulp macrophages
in spleen, alveolar macrophages in lungs and Kupffer cells
in liver [17, 18, 31]. Moreover, equine hyalocytes have a
function in antigen presentation, as evidenced by MHC II expression. MHC II expression usually can be identified in
normal monocyte/macrophages and lymphocytes, and MHC II antigen-bearing cells are considered to be antigen-present-
ing cells [26]. The negative immnoreactive results for CD3, CD20, GFAP and pan-cytokeratin suggested that equine hyalocytes are not differentiated to lymphocytes, glial cells and epithelial cells. S100 protein refers to Ca²⁺ binding pro-
teins that have been found in a variety of cell types, and their localization has been found in potent antigen-presenting cells including dendritic and Langerhans cells [6, 25]. The localization of the S100 protein was not detected in equine hyalocytes of all groups in the present study. In human, the immunohistochemical expression of hyalocytes also provid-
ed strong evidence that hyalocytes belong to the monocyte/
macrophage lineage. However, they antigenically differed from other tissue macrophages in that they express S100 protein and do not express CD68 [19]. The previous study suggested that hyalocytes may originate from a progenitor cell that diverged from the monocyte/macrophage lineage or that expression of these antigens is influenced by differences in the stage of differentiation and/or exposure to modulat-
ing signals from the microenvironment. Equine hyalocytes might have an immunohistochemical phenotype different from that of humans, and that property of those hyalocytes might be influenced by their differentiation and function, and by the different normal ocular microenvironment. Further investigation is needed to clarify the dissimilarity of S100 expression and the delineating functional characterization of equine hyalocytes.

DISCUSSION

CD163 expression of equine hyalocytes was demonstrated in the present study. The expression of these cells indicated that they were resident ocular macrophages in the vitreous cavity. In mouse and rat, hyalocytes also expressed CD163, and the cells were demonstrated as resident ocular macro-
phages [23, 24]. CD163 is a type B crystalline-rich scavenger receptor that is selectively expressed in the cells belonging
to the monocyte/macrophage lineage [9]. The expression is
specifically identified in subpopulations of resident tissue
macrophages in normal tissues, such as interfollicular and
sinus macrophages in lymph nodes, red pulp macrophages
in spleen, alveolar macrophages in lungs and Kupffer cells
in liver [17, 18, 31]. Moreover, equine hyalocytes have a
function in antigen presentation, as evidenced by MHC II expression. MHC II expression usually can be identified in
normal monocyte/macrophages and lymphocytes, and MHC II antigen-bearing cells are considered to be antigen-present-
ing cells [26]. The negative immnoreactive results for CD3, CD20, GFAP and pan-cytokeratin suggested that equine hyalocytes are not differentiated to lymphocytes, glial cells and epithelial cells. S100 protein refers to Ca²⁺ binding pro-
teins that have been found in a variety of cell types, and their localization has been found in potent antigen-presenting cells including dendritic and Langerhans cells [6, 25]. The localization of the S100 protein was not detected in equine hyalocytes of all groups in the present study. In human, the immunohistochemical expression of hyalocytes also provid-
ed strong evidence that hyalocytes belong to the monocyte/
macrophage lineage. However, they antigenically differed from other tissue macrophages in that they express S100 protein and do not express CD68 [19]. The previous study suggested that hyalocytes may originate from a progenitor cell that diverged from the monocyte/macrophage lineage or that expression of these antigens is influenced by differences in the stage of differentiation and/or exposure to modulat-
ing signals from the microenvironment. Equine hyalocytes might have an immunohistochemical phenotype different from that of humans, and that property of those hyalocytes might be influenced by their differentiation and function, and by the different normal ocular microenvironment. Further investigation is needed to clarify the dissimilarity of S100 expression and the delineating functional characterization of equine hyalocytes.

By the histological examination of G2, G3 and G4, equine hyalocytes have been characterized as pleomorphic cells and have been further classified into 4 types in the present study, Hyalocytes in other animals have been known as pleo-

morphic cells in vivo and in vitro [1, 3, 12, 22, 27, 28]. The cell considered to be a type of macrophage in a past study resembled the type-1 cell in the present study, and the type-2 cell was considered to be a subtype of a macrophage [27]. In addition, a fibrocyte type was equivalent to type-3, and the type-4 cell was equivalent to the cell which was a subtype of a fibrocyte type [27]. The morphological differences of hyalocytes were described to reflect the different functions of the cells. Macrophage type was considered to consist of secretory cells responsible for production of hyaluronic acid [13], and fibrocyte type was considered to be the cell responsible for production of vitreous collagen [21]. How-
ever, the evidence of morphological differences in relation to hyalocyte function has not been adequately reproduced. In the present study, the emergence of type-2 and type-4 hya-
locytes tended to be increased in adult horses, and therefore, the morphological changes during process formation were responsible for cell maturity or cell activation. In human and murine, hyalocytes have presented a myofibroblast ap-
pearance, which is potentially relevant to the pathogenesis of cicatricial contractions in proliferative vitreoretinopathies
[14–16]. The present study expounded on the histomorpho-
logical variation of hyalocytes in normal equine eyes. These findings regarding equine hyalocytes may be helpful for the identification of morphological changes which are relevant to the ocular pathological conditions of horses.

In the present study, the site of equine hyalocyte distri-
bution was demonstrated and characterized. The vitreous
cortex of ciliary body was found to be a restricted site of
distribution, and hyalocytes were not located in the vitre-
ous cortex of retina and optic papilla. The free cells found
around ciliary processes were also hyalocytes according to
the findings from the histological and immunohistochemi-
cal examinations. The sites of hyalocyte distribution have
been described previously using various animals [1, 12, 22,
27, 28]. Hyalocytes were located in the ciliary body of all
animal species, but the localization of hyalocytes on retina
was slightly different between animal species, depending on
ocular structure. In guinea pig, which has an avascular retina
similar to horse, hyalocytes were scattered on retina [22].
In chicken, the pectin oculi, which is known as a vascular

---

**Table 2. Primary antibodies used in the present study for immunohistochemical examinations**

<table>
<thead>
<tr>
<th>Primary antibody</th>
<th>Type</th>
<th>Pretreatment</th>
<th>Dilution</th>
<th>Clone</th>
<th>Source</th>
</tr>
</thead>
<tbody>
<tr>
<td>MHC II</td>
<td>MM</td>
<td>Autoclave</td>
<td>1:50</td>
<td>TAL.1B5</td>
<td>Dako</td>
</tr>
<tr>
<td>CD163</td>
<td>MM</td>
<td>Proteinase K</td>
<td>1:50</td>
<td>AM-3K</td>
<td>Trans Genic Inc.</td>
</tr>
<tr>
<td>CD3</td>
<td>RP</td>
<td>Microwave</td>
<td>1:5</td>
<td>—</td>
<td>Nichirei</td>
</tr>
<tr>
<td>CD20</td>
<td>RP</td>
<td>Not done</td>
<td>1:200</td>
<td>—</td>
<td>Thermo Scientific</td>
</tr>
<tr>
<td>GFAP</td>
<td>RP</td>
<td>Microwave</td>
<td>1:100</td>
<td>—</td>
<td>Dako</td>
</tr>
<tr>
<td>Pan-cytokeratin</td>
<td>MM</td>
<td>Autoclave</td>
<td>RU</td>
<td>AE1/AE3</td>
<td>Nichirei</td>
</tr>
<tr>
<td>S100</td>
<td>RP</td>
<td>Microwave</td>
<td>1:300</td>
<td>—</td>
<td>Dako</td>
</tr>
</tbody>
</table>

MM: mouse monoclonal, RP: rabbit polyclonal, RU: ready to use.
structure situated over the head of the optic nerve in this species, was found to be the greatest site of hyalocyte distribution, although they were not distributed on the avascular retina [28]. In human, experimental rodents, and small and large animals, which have blood vessels throughout retina, hyalocytes were found on the retina. Especially in human, cow and cattle eyes, hyalocytes were accumulated linearly along the underlying blood vessels of the retina [1, 27]. In rabbit, hyalocytes have been detected inside the posterior chamber and also have been found on retina [12]. The free cells around the ciliary process were also found in rabbit and guinea pig, as they were in horses [12, 22]. In addition to finding of the site of distribution, the greatest distribution of equine hyalocytes was found at ciliary process. However, pars plana was the site with the greatest number of hyalocytes distributed in calves and cow [27], and has been found inside the vitreous in rabbit [12]. In horses, the results concerning distribution of hyalocytes indicate that ciliary process might play a key role in immigration of equine hyalocytes entering the vitreous via ciliary processes and/or where hyalocytes introduce antigen to the stromal cells.

The finding of equine hyalocyte distributions in G2 suggested that there was immigration of hyalocytes to vitreous cavity before birth. Their site of distribution in G2 was found to be similar to the site of distribution in G3 and G4. In addition, the immunohistochemical expression of hyalocytes in G2 indicated that hyalocytes in G2 had already been differentiated. A significant increase in the number of hyalocytes was detected between fetuses in late gestation and adult horses, and there was no significance between groups of adult horses. This result also indicated that the term from late gestation to after the birth is the time of highest activity for immigration to the vitreous. In a recent study with murine hyalocytes, an age-related increase in the number of subretinal hyalocytes had been described [4, 29, 30]. However, our data did not provide an age-related increase in the number of equine hyalocytes. The factor that brings about an increased number of hyalocytes in normal eyes might vary depending on different ocular microenvironments and/or intraocular physiological conditions.
The present study demonstrated the presence of hyalocytes in the equine vitreous cortex along with their immunohistochemical and histomorphological features and distributions, including the equine fetus. Hyalocytes in other animals including human have been studied under natural and experimental ocular diseases. In human and mouse, hyalocytes participated in proliferative vitreous diseases with myofibroblast appearance and transformation [14–16]. Under experimental ocular diseases, hyalocytes played a role as antigen-presenting cells in ocular immune diseases, responded to systemic LPS exposure and hyperglycemia with increasing their density and productive activation of VEGF in mouse, respectively [14, 24, 29], and represented reactive activations, photocoagulation and vitreous hemorrhage in rabbit [10, 11]. In chicken, hyalocytes exhibited increasing phagocytic activity in maintaining vitreous transparency through the clearance of cellular and extracellular debris, which was provoked by injection of bacteria [28]. Even though there have been many studies of pathological and immunological ocular conditions in domestic and companion animals, hyalocytes in relation to ocular pathological conditions have not been studied. In horses, a variety of ocular diseases have also been reported, but functional contributions of hyalocytes for ocular diseases are still unclear. However, the presence of equine hyalocytes, made evident from MHC II and CD163 expression, is considered to have great potential to play an important role in treating pathological conditions, such as degenerative, proliferative and inflammatory origin, especially in the vitreous cavity. Equine hyalocytes might contribute to progression of the equine ocular pathological conditions as an antigen-presenting cell, and they might participate to those conditions through a phagocytic activity and productions of various physiological active substances. Further investigation is necessary to elucidate participation and actual functions of equine hyalocytes in particular individual ocular pathological condition.

REFERENCES


