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Malignant catarrhal fever-like lesions associated with ovine herpesvirus-2 infection in three 1 2 goats 3 Dr. Björn Jacobsen^a, 4 Dr. Kerstin Thies^b 5 Dr. Alexandra von Altrock^b 6 Christine Förster^c 7 Dr. Matthias König^c 8 Prof. Dr. Wolfgang Baumgärtner^a 9 10 ^aDepartment of Pathology, University of Veterinary Medicine Hannover, Buenteweg 17, D-11 12 30559 Hannover, Germany ^bClinic for Pigs, Small Ruminants, Forensic Medicine and Ambulatory Service, University of 13 Veterinary Medicine Hannover, Bischofsholer Damm 15, D-30173 Hannover, Germany 14 ^cInstitute of Virology, Faculty of Veterinary Medicine, Justus-Liebig-University Giessen, 15 Frankfurter Strasse 107, D-35392 Giessen, Germany 16 17 Corresponding author: 18 19 Prof. Dr. Wolfgang Baumgärtner, Ph.D. 20 Department of Pathology University of Veterinary Medicine Hannover 21 22 Bünteweg 17 23 D-30559 Hannover 24 Tel.: ++49-511-9538621 ++49-511-9538675 25 Fax: Email: wolfgang.baumgaertner@tiho-hannover.de 26

- 27 Abstract
- 28 This is the first description of malignant catarrhal fever-like lesions associated with
- ovine herpesvirus-2 (OvHV-2) infection in goats with central nervous symptoms. The
- diagnosis was based on typical histological lesions characterized by systemic
- 31 lymphohistiocytic and fibrinoid vasculitis and confirmed by polymerase chain reaction and
- 32 subsequent phylogenetic analysis of the detected OvHV-2 sequences.

- 33 Keywords
- Ovine herpesvirus-2, Goats, Vasculitis, Malignant catarrhal fever

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36	Malignant catarrhal fever (MCF), a fatal lymphoproliferative disease of ruminant
37	species including domestic cattle and wild living ruminants, is caused by a group of closely
38	related gammaherpesviruses collectively referred to as malignant catarrhal fever viruses
39	(Callan and Van Metre, 2004). Epidemiologically, two primary forms of MCF in cattle have
40	been described with wildebeest-derived (WD)-MCF, caused by alcelaphine herpesvirus-1
41	(AlHV-1), primarily seen in Africa and the sheep-associated-MCF (SA-MCF) due to ovine
42	herpesvirus-2 (OvHV-2) in North America, Europe and Australia (Callan and Van Metre,
43	2004). In goats another gammaherpesvirus, termed caprine herpesvirus-2 (CpHV-2) was
44	identified (Li et al., 2001) causing MCF in white-tailed deer (Li et al., 2003) and in sika deer
45	(Keel et al., 2003) in the USA. MCF-like lesions or disease in goats have not been reported so
46	far. Interestingly, various ruminant species are susceptible to infection by MCF-viruses, but
47	only few develop characteristic lesions. This is also the case for goats, which can be infected
48	inapparently by OvHV-2 and CpHV-2 (Callan and Van Metre, 2004, Keel et al., 2003).
49	Clinical signs similar to MCF have been observed in sheep under natural conditions, but
50	the conjunction with OvHV-2 was not confirmed by laboratory tests (Rae, 1994).
51	Experimentally, MCF-like disease was induced in sheep after aerosol inoculation with OvHV-
52	2 (Li et al., 2005). MCF was also described in Barbary sheep (Yeruham et al., 2004) and pigs
53	(Loken et al., 1998) identifying OvHV-2 by PCR as the presumable causative agent.
54	Here we describe MCF-like lesions in goats with OvHV-2-infection as confirmed by
55	PCR and nucleic acid sequencing.

Results and discussion

57	Three female goats, two German Improved Fawn breed (No. 1, 3.5 years old; No. 2, 7
58	years old) and one dwarf goat (No. 3, 5 month old) originated from two different flocks
59	consisting of three goats and five sheep, and three dwarf goats, respectively. All three
60	animals showed fever and central nervous disorders including ataxia, tonic-clonic spasm,
61	disorientation and tremor of the head. Animal No. 3 displayed diarrhea and cloudy corneas in
62	both eyes. Goat No. 1 was euthanized by the local veterinary surgeon after 7 days of clinical
63	disease, goat No. 2 and No. 3 died after 22 and 23 days of illness, respectively. At necropsy,
64	the animals showed enlargement of lymph nodes, spleen and liver, pulmonary congestion and
65	multifocally distributed tiny white spots in liver and kidney. Goat No. 3 displayed, in
66	addition, an erosive, ulcerative esophagitis. Histopathology on hematoxylin and eosin stained
67	sections revealed a lymphohistiocytic vasculitis with fibrinoid necrosis of vessel walls in
68	several organs including lymph nodes, liver, kidney, brain, retina, and rete mirabile (Figure
69	1). Additionally, a granulomatous pneumonia due to parasites (No. 1), lymphohistiocytic
70	meningitis (No. 1), pericholangitis (No. 2) and a multifocal ulcerative lymphohistiocytic
71	esophagitis (No. 3) were observed.
72	Immunhistochemically, antigen for Borna disease virus, porcine herpesvirus-1,
73	Toxoplasma gondii and Neospora caninum were not detectable. By immunofluorescence
74	microscopy antigens of rabies and pestiviruses were not found.
75	Samples from different organs (goat No. 1: brain, kidney; goat No. 2: brain,
76	cornea, lymph node, blood; goat No. 3: brain, mesenteric lymph node, spleen) were used
77	for two different PCR set ups. Primers targeting the tegument region of OvHV-2 were taken
78	from the literature (Baxter et al., 1993). For the glycoprotein B region of OvHV-2 and CpHV-
79	2 primers were designed using the primer express software (ABI, Langen, Germany). The
80	primer pair designated GlycoF02 (5'TCYGCCAAGCGMATCG3') and GlycoR01
81	(5'ACCGTGCTCATGCGCTT3') amplified a 424bp fragment. Upon gel electrophoresis all

samples showed DNA bands at the expected size. Positive controls for OvHV-2
originated from the brain of a cow that died after showing clinical signs of MCF and
peripheral blood lymphocytes from an inapparently infected goat for CpHV-2. Obtained
reaction products were cloned and sequenced. Comparison to nucleotide sequences of the
glycoprotein B region of other herpesviruses retrieved from the gene bank revealed
similarities of 99.04 - 99.68% for OvHV-2 and 83.93% for CpHV-2, respectively. Based on the
DNA sequences of the glycoprotein B fragment including several other members of the
gammaherpesvirinae pair wise genetic distances were calculated using the Kimura 2-
parameter method (Kimura, 1980) and a phylogenetic tree was constructed applying the
neighbour-joining method (Felsenstein, 1985) from the PHYLIP software package
(University of Washington, Seattle, USA) (Figure 2). Phylogenetic analysis based on
amplicons from the tegument region led to the same results (data not shown).
Clinically, these animals showed central nervous disorders, increased body temperature,
pleocytosis and diarrhea. Similar signs have been described for cattle (Callan and Van Metre,
2004). Frequent clinical signs in bovine include persistent high fever and lymph node
enlargement, whereas the intestinal form characterized by diarrhea is less often observed in
this species (Callan and Van Metre, 2004). Neurological signs in bovine are less frequent and
typically appear in the terminal stages of the disease. Affected animals are lethargic and
appear obtunded (Callan and Van Metre, 2004). In the cerebrospinal fluid there is a marked
pleocytosis (Plowright, 1990). Though animal No. 3 displayed a cloudy cornea, characteristic
lesions of the "head-and-eye"-form described for cattle were not seen in these goats.
Macroscopically, all three goats showed enlargement of lymph nodes and visceral organs with
tiny white spots in liver and kidney. One animal displayed ulcerations and erosions of the
esophagus. Similar findings have been reported for MCF in cattle (Plowright, 1990).
Histopathologically, the animals showed lymphohistiocytic vasculitis with fibrinoid necrosis
in several organs. In cattle this regarded as almost nathognomonic for MCF (Callan and Van

108 Metre, 2004). Vasculitis in bovines, similar to the present cases, is segmental and irregular in 109 distribution, most readily seen in medium-sized arteries invariably accompanied by perivascular and intramural infiltration of mononuclear cells (Plowright, 1990). Frequently, 110 111 there is fibrinoid degeneration of infiltrated mural tissues and smooth muscle elements of 112 vessel walls. Vascular lesions appear to have predilection sites including kidney, brain and the 113 rete mirabile of the hypophysis (Plowright, 1990). Until now, such disease pattern has not 114 been described in goats. Though the pathology of MCF is well documented, the pathogenesis still is poorly 115 116 understood. Studies suggest a lymphoproliferative disorder involving the dysregulation of Tlymphocytes (Callan and Van Metre, 2004). In addition, there is evidence that MCF presents 117 an immunopathological process (Plowright, 1990). The essential defect in MCF pathogenesis 118 119 appears to be an immune deregulation attributable to a dysfunction of Natural Killer cells and uncontrolled proliferation of lymphoblastoid elements in many tissues (Plowright, 1990). 120 121 Whether MCF-lesions are due to a type III or Arthus-like hypersensitivity reaction is still 122 discussed controversially (Plowright, 1990). Most episodes of spontaneous SA-MCF in cattle 123 appear to be a sequel of close contact to infected sheep actively shedding the agent (Metzler, 1991). Goat No. 1 and No. 2 were kept together with sheep. Though, goat No. 3 was kept 124 125 together with sheep too, direct contact has been ruled out by the owner. However, there is 126 evidence that OvHV-2 transmission can occur over distances (Callan and Van Metre, 2004). Confirmatory diagnosis was based on the histological demonstration of a generalized 127 lymphohistiocytic vasculitis in multiple organs, including the brain. Since members of the 128 129 gammaherpesvirinae in general cannot be propagated easily in cell culture, PCR represents the method of choice to facilitate a direct diagnosis of viral infection. **Infections of goats** 130 with both OvHV-2 and CpHV-2 have been reported (Li et al., 2001, Metzler, 1991). 131 132 However, data with regard to the prevalence of both agents in the goat population are 133 limited. On the other hand OvHV-2 and CpHV-2 are well known as agents of MCF in

et al., 2003, Yeruham et al., 2004, Plowright, 1990), however, MCF or MCF-like lesions have not been described in goats so far. Unfortunately, data of the OvHV-2 status of the sheep and unaffected goats from both flocks are not available. Experimental infections demonstrated the crucial role of the viral dose with regard to disease development in sheep and in cattle (Li et al., 2005, Taus et al., 2006). The latter studies suggested that a high viral burden correlates with the development of clinical MCF and histological lesions (Taus et al., 2006, Li et al., 2005). If a similar pathogenesis applies to OvHV-2 infection in goats remains to be investigated in further studies.

This is the first known description of MCF-like lesions in goats associated with an infection with OvHV-2. Diagnosis of OvHV-2 was established by PCR and confirmed by nucleotide sequencing. Phylogenetic analysis, based on two different regions of the genome, identified the viruses detected in three infected goats as OvHV-2 that is clearly separated from known caprine herpesviruses. Although the etiological role of the viruses found independently in goats with signs of MCF was not formally established, it is likely that OvHV-2 was

responsible for disease induction and played an essential role in the pathogenesis of the

lesions. However, further studies are required to establish a causal relationship between

the observed clinic, lesions and OvHV-2 infection.

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190	Figure capuons
191	Figure 1:
192	Lymphohistiocytic vasculitis (star) with fibrinoid necrosis (arrow) in a medium-sized artery,
193	kidney, hematoxylin and eosin. Bar 50μm.
194	
195	Figure 2:
196	Phylogenetic analysis of the glycoprotein B sequence (315bp): The branch lengths are
197	proportionate to relative genetic distances. Statistical significance of the branches was
198	calculated by boot strap analysis using 1000 replicates. Boot strap values are indicated for
199	main branches. According to the phylogenetic analysis the newly identified sequences from
200	the three goats with clinical symptoms of malignant catarrhal fever (Goat No. 1, No. 2 and
201	No. 3) group together with ovine herpesvirus-2 (OvHV-2 I-V [AF327833, AF385442,
202	AF385439, AF385440, AF385441]; 99,04-99,68% similarity) and clearly separate from
203	caprine herpesvirus-2 (CpHV-2 [AF283477]; 83,93% similarity), alcelaphine herpesvirus-1
204	(AlHV-1 [AF005370]; 79,01-79,89% similarity) and other tested members of the
205	Gammaherpesvirinae (alcelaphine herpesvirus-2 (AlHV-2 [AY237369]), bovine
206	lymphotropic herpesvirus (BLHV I and II [AF327832, AY237373]), type 1 and 2 ruminant
207	rhadinovirus (Type 1 RRhV [AY237371]; Type 2 RRhV I and II [AY237367, AY237368]),
208	and malignant catarrhal fever virus in white-tailed deer (WTD-MCF [AY237370]).



