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Bovine Viral Diarrhea-an Emerging Disease in Camelids a Review

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ABSTRACT

Bovine Viral Diarrhea (BVD) is an emerging disease in both New World Camelids (NWCs) and Old World Camelids (OWCs). The virus has been isolated from NWCs particularly in alpacas and dromedaries, but there are no reports of BVD in Bactrians. BVD is an important infectious disease. Both sub-genotypes 1a, 1b and genotype 2 have been isolated from NWCs but the ncp BVDV 1b is primarily implicated in cases of BVD in NWCs. A BVD strain unique to camelids has not yet been isolated. In NWCs virtually all infections have been caused by the non-cytopathic (ncp) BVDV, Persistently infected crias have also been detected. Llamas and alpacas demonstrate clinical signs such as ill thrift, diarrhea, respiratory ailments and abortions. As in bovines, identification and elimination of PI animals, has the highest priority to avoid infection of the entire herd. BVD was also observed in dromedaries in Egypt. Both isolates revealed a cytopathic effect (cpe) and so far no ncp virus has been isolated from dromedaries. Also in dromedaries, BVD infections caused intrauterine death, stillbirth, weak calf syndrome with congenital deformities, neonatal respiratory disorders in young dromedary calves and acute hemorrhagic gastroenteritis in adult dromedaries. So far, no PI dromedaries have been described.

Keywords: Bovine Viral Diarrhea Virus (BVDV), Mucosal Disease (MD), Old World Camels (OWCs), New World Camels (NWCs), Persistently Infected (PI)

1. INTRODUCTION

Bovine Viral Diarrhea Virus (BVDV) is responsible for two distinct clinical entities in cattle which are:

- Bovine Viral Diarrhea (BVD) resulting in high morbidity and low mortality
- Mucosal Disease (MD), which is sporadic but regularly fatal

Two genotypes of BVDV are recognized (BVDV-1, BVDV-2) and each genotype has 2 biotypes: noncytopathic (ncp) and cytopathic (cp). The nature of biotypes involved in BVDV infection play a pivotal role in the epidemiology and pathogenesis of the disease not only in cattle but also in camelids. Only ncp strains of BVDV produce Persistently Infected (PI) animals, whereas cp strains are derived by mutation from preexisting ncpBVDV strains and do not produce viremic animals. Only ncp strains are excreted and may infect other animals in a herd. These strains are also able to cross the placental barrier and infect the fetus in the early stages of gestation. Bovine Viral Diarrhea (BVD) and Mucosal Disease (MD) are epidemiologically different diseases in cattle that have different pathogeneses, although both are caused by the same *Pestivirus*. *Pestivirus* infection can occur at any age in postnatal life and may be subclinical or produce a range of clinical conditions including acute diarrhea, acute hemorrhagic syndrome, acute fatal and wasting disease. The *Pestivirus*, widespread in cattle populations worldwide has also been isolated from New World Camels (NWCs) and Old World Camels (OWCs) (Evermann *et al.*, 1993; Mattson, 1994; Hegazy *et al.*, 1998).

Over many years a great amount of research has taken place mainly in NWCs in connection with the *Pestivirus*, which resulted in a number of publications as summarized in **Table 1**. It is now obvious that BVDV can produce cases of diarrhea, ill thrift, reproductive losses, respiratory disease and disseminated disease in NWCs (Kapil *et al.*, 2009). A review article on BVDV infection in NWCs has been recently published by Amstel and Kennedy (2010).



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	NWCs			
Author	Year	Country	Species	Result
Rivera et al.	1987	Peru	Alpaca	serology 11% reactors
Karesh et al.	1998	Argentina	Guanaco	serology all negative
Belknap et al.	2000	USA	Llama	virus isolated FAT negative immunoperoxidase positive SNT negative
Goyal et al.	2002	USA	Alpaca	ncp type 1b isolated, PCR positive
Wentz et al.	2003	USA	Llama	experimental infection no disease
			Alpaca	seroprevalence is low no fetal infection
Carman <i>et al</i> .	2005	USA	Alpaca	Systemic disease abortion, virus isolation, persistent infection, BVDV type 1b
Anonymous	2005	UK	Alpaca	Systemic disease abortion, ncp virus isolation, BVDV type 1b
Anonymous	2005	UK	Alpaca	Ill thrift, diarrhea, type 1b BVDV
Foster <i>et al</i> .	2005	UK	Alpaca	ncp BVDV type 1
Evermann	2006	USA	Llama	1 51
			Alpaca	General article about BVD in llama and alpaca
Henningson et al.	2006	USA	Alpaca	Persistently infected, pathological lesions
Bromage	2006	UK	Alpaca	General article
Anonymous	2006a	UK	Alpaca	PCR positive for BVD type 1, abortion
Anonymous	2006b	UK	Alpaca	Abortion, weak crias, PCR, BVD type 1 positive
Anonymous	2006b	USA	Camelids	General article
Bedenice	2006	USA	NWCs	General article
Evermann et al.	2006	USA	Camelids	General article
Mattson <i>et al</i> .	2006	USA	Alpaca	Persistent infection, BVDV isolated
Mueller and Broadbent	2007	USA UK	Camelids	General article
Probst et al.	2007	European zoos	Camelids	Serology 1.4%
Kelling	2008	USA	Alpaca	Serology 25.4% herds were positive
Byers	2008	USA	Camelids	General article
Bedenice	2008	USA	Alpaca	Persistent infection with BVDV type 1
Danuser et al.	2009	Switzerland	Alpaca	
			Llama	Seroprevalence 4.6%
Shimeld	2009	USA	alpaca	Seroprevalence 20%, persistently infected animals
Kim et al.	2009	USA	alpaca	46 BVD viruses identified by PCR, one genotype 1b BVDV
Topliff et al.	2009	USA	NWCs	20% seroprevalence, 6% PI crias
Mudry <i>et al</i> .	2010	Switzerland	NWCs	Serology 5.8%, 3.6%, no pestiviral RNA found
Johnson et al.	2010	USA	Alpacas	Intranasal infection with BVDV 1b and 2 from bovine and BVDV 1b from alpaca
Byers et al.	2010	USA	Alpacas	Vaccination trial
Byers et al.	2011	USA	Alpacas	Exposure of native alpacas from PI alpacas
Bedenice et al.	2011	USA	Alpacas	35 crias naturally infected with BVDV type 1b
Doyle and Heuschele	1983	USA zoos	Dromedary	Serology 13%
Fahmy	1999a, 1999b, 1999c	Egypt	Dromedary	Experimental infections
Yousif et al.	2004	Egypt	Dromedary	BVD virus type 1 and 2 isolated
Al-Afaleq et al.	2006	Saudi Arabia	Dromedary	Seroprevalence 18%
Taha	2007	UAE	Dromedary	Seroprevalence negative
Wernery et al.	2008	UAE	Dromedary	Seroprevalence in camel dairy 1.6%



1.1. Etiology

Bovine Viral Diarrhea virus (BVDV) is a small RNA virus of the *Flaviviridae*. Together, with the viruses of border disease and classical swine fever virus, it forms the genus *Pestivirus*.

The three viruses are antigenically related. Strains isolated from newborn calves and persistently infected cattle, are generally non-cytopathic (ncpBVDV), while those from tissues of cattle suffering from MD are usually cytopathic (cpBVDV). Today two genotypes of BVDV are recognized: BVDV-1 and BVDV-2 based on the nucleotide sequence of the 5'- untranslated region. BVDV-1 has a worldwide distribution, whereas BVDV-2 is largely restricted to the USA and Canada but has also been isolated in Italy, Holland and the UK (Reed, 2010). Each of the two genotypes has two biotypes, noncytopathic (ncp) and cytopathic (cp) (Peterhans et al., 2010). Genotype 1 isolates are further divided into at least 2 sub-genotyes (1a and 1b). Particularly in alpacas both sub-genotypes 1a, 1b and genotype 2 have been isolated (Amstel and Kennedy, 2010). Non cytopathic BVDV 1b is primarily found in NWCs. A BVDV strain unique to camelids has not yet been identified.

1.2. Epidemiology

Postnatal infection with the virus is acquired by ingestion or inhalation of contaminated material which results in the development of serum neutralizing antibodies. This is usually a clinically unrecognizable infection. On the other hand, with infection of a nonimmune pregnant animal, the virus always crosses the placental barrier and invades the fetus. While the dam seroconverts without showing signs of disease, the fetus is immunotolerant in the early stages of pregnancy. This congenital infection can result in a wide spectrum of abnormalities; fetal death, congenital defects, or a persistent lifelong infection without clinical signs. The outcome is mainly dependent on the stage of fetal development during which time the infection takes place. BVDV is also transmitted in semen, particularly from persistently infected bulls, which shed the virus in their semen for their whole life.

Many years were required before understanding the complexity of infection with the *Pestivirus*, especially the link between BVD and MD in cattle. *Pestivirus* infection has only been recently recognized as a potential cause of serious illness in both NWCs and OWCs (Evermann, 2006), indicating that members of the camelid group are susceptible to infection. In NWCs, *Pestivirus* has been intensively researched in North

America (Byers, 2008; Kim et al., 2009), the UK (Mueller and Broadbent, 2007) as well as in Switzerland (Danuser et al., 2009). BVDV may cause a severe disease in alpacas and llamas, including diarrhea, reproductive loss, wastage and death, posing a significant threat to a herd's health as it does in bovines (Belknap et al., 2000). The emerging BVDV infections have raised significant concerns in the camelid industries in North America and Europe, especially in the UK, Switzerland and Germany that have strong NWC societies. Similar to cattle, there is not only a systemic disease caused by the Pestivirus but also persistent BVDV infection in NWCs may occur, when the fetus is infected during early pregnancy, before it becomes immunocompetent. Field observations have shown that up to 80% of crias born to naïve dams infected during early pregnancy may become Persistently Infected (PI) with the Pestivirus (Bedenice, 2008; Bedenice et al., 2011). At a population level in bovines, the prevalence of PI animals is about 1%. So far, nearly all infections in SACs are caused by the ncpBVDV including the Persistently Infected (PI) crias. Only in one case was a cytopathic biotype documented from a llama foetus (Bedenice, 2006). However, serological investigations by Shimeld (2009) using serum virus neutralization (SN) with NADL (BVDV type 1) and c125 BVDV (BVDV type 2) on 426 alpaca sera, found that 20% of the alpaca, were sero-positive for one or both genotypes in the US.

PI animals are pivotal in the epidemiology of the disease (Byers et al., 2009) because they disseminate not only the ncpBVDV which they harbour, but also the cp BVDV derived by mutation. These PI animals must be eliminated from the herd as soon as possible. From 2004 to 2007 more than 12,000 alpacas in the US were screened by real-time PCR, to identify alpacas persistently infected with BVDV. A total of 46 BVDV isolates were found and analyzed by comparison of nucleotide sequences of 2 viral genome regions. The results showed that unique genotypes of bovine BVDV 1b are maintained in the alpaca population of the US. It is not yet clear why alpacas were predominantly infected with genotype 1b BVDV isolates and how bovine BVDV evolved to infect alpacas, although camelids are susceptible to other genotypes (Kim et al., 2009). It is thought, that BVDV infection of camelids primarily originates from infected cattle via intermingling (Evermann, 2006).

The most likely route of BVDV infection in camelids is via the oronasal mucosa most probably after inhalation of viral particles which are present in body fluids of infected animals (Byers *et al.*, 2011).



In the US several other researchers have isolated the 1b BVDV (Carman *et al.*, 2005) but the same genotype was also found in alpacas in the UK (Anonymous, 2005; Fahmy, 1999c) and Switzerland (Danuser *et al.*, 2009). Mueller and Broadbent (2007) reported that the oldest surviving PI alpaca survived 30 months. In bovines, more than 50% of PI animals die before they reach the age of one year. PI bovines are mostly clinically healthy and excrete large quantities of BVDV, which is antigenically identical to the ncpBVDV present in the animal (Lefèvre *et al.*, 2011). A similar pathogenesis can be expected to occur in NWCs but has not been described in OWCs. A report by Carman *et al.* (2005) indicates a BVDV infection, *in utero*, results in immunotolerant crias.

It is known that both llamas and alpacas are susceptible to BVDV infection and that in some cases, the animals demonstrate clinical signs such as diarrhea, ill thrift, respiratory ailment and abortion. BVD can be severely exacerbated in connection with a bovine tuberculosis (bTb) infection, leading to fatalities (Cobb and Cobb, 2010). It is also assumed that male camelids could, like bulls, persistently be infected.

Several researchers have also stressed that BVD is uncommon in SACs, despite the fact that 25% of the alpaca herds can be serologically positive (Kelling, 2008) with a seroprevalence of 20% (Shimeld, 2009). The Alpaca Research Foundation of the US, believes that the incidence of clinical BVD in the US, is only 0.05% (Evermann et al., 2006; Shimeld, 2009). Topliff et al. (2009) reported in a recent nationwide survey of 63 herds, a prevalence of 25% seropositive crias, of which 6% had evidence of BVDV PI crias. Belknap et al. (2000) found a serological BVDV incidence in NWCs of 4.4% and Evermann (2006) as high as 53%. In a comparative epidemiological study, conducted by Mudry et al. (2010) in Switzerland on llamas and alpacas in 2000 and 2008, it was found that Pestivirus infection is a negligible risk for the BVDV eradication programme. seroprevalence and the pestiviral The RNA seroprevalence was low, with 5.8% versus 0% in 2008 and 3.6% versus 0% in 2000. Byers et al. (2011) reported only mild sign in BVDV naive alpacas after they were exposed to two PI alpacas which shed BVDV type 1b virus in most of their body fluids. Viremia was detected in the transient infected alpacas but viral shedding during the acute phase did not take place and antibodies appeared to be protective upon re-exposure to the type 1b virus.

Two *Pestivirus* isolated by Hegazy *et al.* (1995) in Egypt, from adult dromedaries with diarrhoea and from dromedary calves born with congenital defects, were

genotyped by Yousif *et al.* (2004). Genotyping revealed that isolate Giza4 and isolate Giza7 belong to BVDV-1 and BVDV-2, respectively.

1.3. Clinical Signs and Pathology

Serological studies indicate that NWCs and OWCs are susceptible to infection with the BVDV. The results of serological studies identifying BVDV antibodies in the dromedary have appeared, but no reports have been from Bactrian camels. In dromedaries, found seroprevalences between 2 and 50% have been reported. BVDV antibodies have appeared in dromedaries from Tunisia, with 3.9% positive cases (Burgemeister et al., 1975), from Oman, 6.7% positive cases (Hedger et al., 1980), from Sudan, 15.5 and 15.7% positive cases (Bornstein and Musa, 1987; Bornstein et al., 1989) and from Somalia with 3.4% positive cases (Bornstein, 1988). Bohrmann et al. (1988) did not identify any antibodies to BVDV in Djibouti using the serum neutralization test. Wernery and Wernery (1990) explained the higher incidence of BVD in UAE breeding camels (9.2%) when compared to racing dromedaries (3.6%), with their larger breeding herds and closer contact with cattle herds. In a later survey (CVRL, 1998), these findings were confirmed using an antibody ELISA. The incidence of BVDV antibodies in 552 camels tested was, 0.5% in racing camels and 6.4% in breeding camels. The presence of neutralizing antibodies to BVDV was 11% in Egypt, with a peak of 23% in one area (Hegazy et al., 1993). In another Egyptian survey, Tantawi et al. (1994), detected 4.3% BVDV positive dromedaries and Zaghana (1998) found that camels from Egypt, exhibited an even higher prevalence (52.5%) of neutralizing antibodies to BVDV.

Doyle and Heuschele (1983) examined 24 sera with the SNT from dromedaries kept in American zoos, of which 13% had antibodies using the Singer BVDV strain. Newer seroepidemiological studies from Saudia Arabia and the United Arab Emirates showed a high seroprevalence with 18% (Al-Afaleq et al., 2006) in Saudia Arabia and a low seroprevalence in the UAE, with 1.6% in dairy dromedaries (Wernery et al., 2008) and no reactors in 812 sera by Taha (2007). As in OWCs, many BVD serological surveys have been conducted in NWCs, as can be seen from Table 1. In a serological survey conducted in Peru, involving 117 alpacas that grazed with cattle and sheep, the prevalence of antibodies to BVDV was 11% (Rivera et al., 1987) and Picton (1993) reported a prevalence of 4.4% in 270 llamas from Oregon in the USA. A study by Puntel et al. (1999) found 2.05% (8/390) reactors to the BVDV in



llamas, from nine farms located in three different provinces in Argentina.

Cattle suffering from BVD and MD show lesions in the alimentary tract. The pathological changes in MD are much more severe than in BVD, the MD lesions are often found only in the upper alimentary tract. In both BVD and MD, pathological changes consist mainly of erosions and ulcers of varying severity. These pathological lesions have not been described in NWCs, but PI alpacas revealed multifocal hepatic necrosis and bronchopneumonia, as well as severe thymicatrophy, as seen in bovine calves (Henningson et al., 2006). BVD viral antigen was widely distributed in the alpaca tissues mostly prominent in nerves, Tunica media and adventitia of vessels and macrophages in the gastrointestinal submucosa, urinary bladder submucosa, lungs and cerebrum, indicating a viremia. The authors concluded that PI alpacas thymicatrophy may indicate hosts defense deficits, similar to PI bovine calves.

Long-term clinicopathological characteristics of alpacas naturally infected with BVDV type 1b were reported from Bedenice *et al.* (2011). Their investigation involved 35 crias which were naturally exposed to BVDV. Chronically infected and PI crias had developed a significantly lower birth weight, decreased growth rates, anemia and a monocytosis compared with control animals. BVDV type 1b infection during early pregnancy resulted in a high incidence of PI crias which developed chronic wasting, diarrhea and severe respiratory distress.

Three groups of BVDV naïve alpacas containing six animals each where intra nasally infected with BVDV 1b of bovine origin (group1), BVDV2 of bovine origin (group2) and BVDV 1b of alpaca origin (group3) to study clinical signs, viremia, seroconversion and hematological changes (Johnson *et al.*, 2010). After infection all three genotypes induced nasal BVDV shedding, viremia and sero- conversion but there were differences in the median onset.

It is reported that alpacas develop a milder disease following BVDV infection compared to bovines which was investigated by Samson *et al.* (2011). The authors found that the limited permissiveness of alpaca cells to BVDV compared to bovine cells may be the reason for this phenomenon.

BVD infections have been described in dromedary calves from Egypt (Hegazy *et al.*, 1998) causing intrauterine death, stillbirths and weak calf syndrome with congenital deformities, neonatal respiratory distress syndrome in young dromedary calves and acute hemorrhagic gastroenteritis in adult dromedaries. BVDV was isolated from lymphoid tissues, spleen, brain and kidney on Bovine Kidney Cells (BKC) causing a cytopathic (cpe) effect. The virus was also observed by immunofluorescence in different organs. The virus was then later genotyped by Yousif *et al.* (2004) and named Giza 4 and Giza 7, belonging to BVDV type 1 and BVDV type 2. No ncpBVDV was isolated and no reports exist that PI animals have been detected in dromedaries. Extensive studies have been carried out by Fahmy (1999a; 1999b; 19999c) on pregnant goats infected with the Egyptian isolates and a bovine BVD virus. These experimental trials included:

- Studies on the reproduction performance
- Studies on the effect of the fetus and newborn kids
- Clinicopathological studies

The inoculation of pregnant does at day 65 of gestation, with the bovine NADL BVDV, resulted in early abortions in 60% of inoculated does, whereas the cpe camel strain induced late abortions in 25%. The aborted fetuses caused by both viruses were small in size, severely autolysed and had been obviously dead in uterus for a long period of time. The inoculated does revealed severe leukopenia and lymphopenia during the first 7 days p.i. with both strains. Lymphopenia lasted for 28days. The remaining does gave birth to healthy kids with weight comparable to control kids.

Hegazy *et al.* (1995) state that the main cause of abortion in dromedaries is caused by the BVDV, which can reach 50% in some herds. This statement is very controversial, as no abortions storm have been described in other countries.

In the UAE for example, adult dromedaries and calves that have died of other causes, are routinely virologically screened, including the fluorescence test for the presence of the BVDV. So far the results have always been negative (Wernery *et al.*, 1992).

1.4. Diagnosis

Diagnosis of BVD and MD requires laboratory support in the form of virus isolation, virus antigen detection and serum antibody determination. Skin biopsies are the tissues of choice for the diagnosis of BVDV, using immunohistological techniques and are always positive in persistently infected animals (Braun *et al.*, 1999). This method should also be applied in the diagnosis of this disease in camelids (Evermann *et al.*, 1993).

Goyal *et al.* (2002) stressed multiple test systems should be applied for arriving a proper diagnosis because their immunhistochemical examination of



various tissues were negative although a ncp type 1b was isolated from the tissues.

Diagnosis of BVD in cattle must be carried out at two levels. Firstly, the serological status of the herd must be determined either by serum neutralization tests or by ELISAs. Competitive ELISAs have been developed and are ideal for serodiagnosis of camelids. If no reactor is found in a herd, the herd is considered free of BVD. On the other hand, if one or several animals are detected with BVDV antibodies, it is essential to search for PI animals. The presence of PI animals may gradually contaminate the entire herd and therefore, the identification and elimination has the highest priority. For the detection of PI animals, capture ELISAs and PCR assays are nowadays used. PCR assays can also differentiate between strains of type 1 and type 2. Antibody ELISAs will not detect PI animals that are immunotolerant.

1.5. Treatment and Prevention

Knowledge and technical tools to control BVD are readily available and national control schemes in several European countries are now in place (Brownlie *et al.*, 2000).

Economic losses caused by BVD/MD mainly arise from prenatal infections. It is therefore essential to remove all persistently infected animals and to vaccinate heifers prior to first breeding (Thiel et al., 1999). The main risk is the re-introduction of infection into a herd, through a serologically positive pregnant dam carrying an infected fetus, which will always give birth to infected animal. Since it is known that BVDV also causes abortions in camels, it may be necessary to adopt control and vaccination strategies similar to those carried out in cattle. Live and inactivated vaccines have been widely used in several countries. Live vaccines are not recommended in camelids, because a variety of adverse effects, have been observed using live BVDV vaccines in cattle. However, Byers et al. (2010) did not observe any adverse effects when nonpregnant female alpacas were vaccinated with a modified live BVDV vaccine and challenged 25 days post immunization by nasal and ocular inoculation with BVDV type 1b strain. The modified-live BVDV type1 (Singer strain and BVDV type 2 (srain 125c) vaccine (Vista 3SC vaccine, Intervet/Shering Plough Animal Health, Millsboro, DE USA) was administered subcutaneously at a dose of 2 ml according to the recommendation for use in cattle. Results of this vaccination experiment showed that the type 1b BVDV challenge strain which originated from a persistently infected alpaca was not

detected from the vaccinated alpacas but from the two unvaccinated controls.

However, inactivated vaccines are safer and can provide good protection. New developments have led to the production of vaccines from ncpBVDV strains. It has been shown that NWCs seroconverted after a regimen of three vaccinations using an inactivatedvirus preparation (Mattson, 1994).

2. CONCLUSION

Over the last years our knowledge of BVD in camelids has increased and it is obvious that both NWCs and OWCs can contract the disease. However, extensive studies are necessary to elucidate the entire disease pattern in this animal species, as with bovines, through extensive field observations and laboratory studies. Investigations in bovines, have led to a new understanding of the complex epidemiology and pathogenesis of BVD and MD and one can hope that this will also be the case for the camelid family.

3. ACKNOWLEDGEMENT

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