Monitoring sheep and *Culicoides* midges in Montana for evidence of Bunyamwera serogroup virus infection

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**ABSTRACT**

**Introduction:** A serological and entomological investigation was performed to monitor for potential Bunyamwera (BUN) serogroup virus activity in Montana.

**Results:** To facilitate the serological investigation, sera were collected from 104 sheep in 2013 and 2014 and assayed by plaque reduction neutralization test using all six BUN serogroup viruses known to occur in the United States: Cache Valley virus (CVV), Lokern virus (LOKV), Main Drain virus (MDV), Northway virus, Potosi virus and Tensaw virus. BUN serogroup virus-specific antibodies were detected in 41 (39%) sheep. Of these, three were seropositive for MDV, one was seropositive for CVV, one was seropositive for LOKV and 36 had antibodies to an undetermined BUN serogroup virus. Additionally, 30,606 Culicoides sonorisensis were collected in 2013 using Centers for Disease Control and Prevention (CDC) light traps and assayed for cytopathic virus by virus isolation in African Green Monkey kidney (Vero) cells. All midges were negative. Almost one-third of the midges were further tested by reverse transcription-polymerase chain reaction using BUN serogroup virus-reactive primers and all were negative.

**Conclusions:** We provide evidence of BUN serogroup virus infection in sheep but not C. sonorisensis in Montana in 2013-2014. This study also provides the first evidence of CVV, MDV and LOKV activity in Montana.

**INTRODUCTION**

All viruses in the Bunyamwera (BUN) serogroup (genus *Orthobunyavirus*, family *Bunyaviridae*) are maintained in transmission cycles involving haematophagous arthropods and vertebrate hosts (Schmaljohn and Nichol 2007). Six BUN serogroup viruses occur in the USA: Cache Valley virus (CVV), Lokern virus (LOKV), Main Drain virus (MDV), Northway virus (NORV), Potosi virus (POTV) and Tensaw virus (TENV) (Calisher and others 1986, Francy and others 1990). CVV infections in sheep can result in embryonic and fetal death, stillbirths and congenital malformations (Edwards and others 1989, Chung and others 1990a, b). MDV induces severe musculoskeletal and nervous system malformations in ovine fetuses infected in utero by injection into the amniotic vesicle and is a cause of equine encephalomyelitis but has not been associated with naturally occurring disease in sheep (Emmons and others 1983, Edwards and others 1997). LOKV, POTV, NORV and TENV are not recognised ovine pathogens; however, their ability to cause disease has not been widely investigated. Because most veterinary diagnostic laboratories do not routinely test for BUN serogroup viruses, information on their ability to cause disease and data on their true disease incidence and seroprevalence in livestock are limited.

Most viruses in the BUN serogroup are transmitted primarily by mosquitoes (Calisher and others 1986). Several species of biting midges in the genus *Culicoides* also play an important role in BUN serogroup virus transmission. LOKV and MDV have been repeatedly isolated from *Culicoides* species, and *Culicoides variipennis* is a competent vector of MDV (Mellor and others 1974, Calisher and others 1986). *Culicoides* species are also important vectors of several medically and veterinary important orthobunyaviruses in the Simbu serogroup (e.g. Akabane virus, Oropouche virus and Schmallenberg virus (SBV)) (Mellor and others 2000, Rasmussen and others 2012, Veronesi and others 2013). For instance, field studies in Denmark (Rasmussen and others 2012), Italy (Goffredo and others 2013) and Belgium (De Regge and others 2012) have implicated biting midges of the *C. obsoletus* group as relevant vectors of SBV and *Culicoides sonorensis* was demonstrated to be a suitable model vector species for SBV transmission (Veronesi and others 2013). *C. sonorenensis* is an abundant species in Montana (Johnson, unpublished data), produced in the wet,
manure-contaminated soil surrounding stock ponds and other lentic water sources on farms and ranches.

There is no recent information on the seroprevalence of BUN serogroup viruses in livestock in Montana, nor are there any recent data of the significance of Culicoides species in BUN serogroup virus transmission in this region. Therefore, the overall goal of this study was to perform a serological and entomological investigation to determine the seroprevalence of BUN serogroup viruses in sheep and the prevalence of these viruses in C. sonorensis temporally and spatially associated with sheep in Montana.

MATERIALS AND METHODS

Sera collections

Sera were collected from sheep from a ranch in south central Montana in 2013 and a ranch in eastern Montana in 2014 (Fig 1). Several ranchers were contacted in the study area where Culicoides were collected regarding participation in the study. Some declined and others were unable to participate because of other commitments.

Plaque reduction neutralisation tests

Plaque reduction neutralisation tests (PRNTs) were performed using CVV (strain CVV-478), LOKV (strain FMS 4352), MDV (strain BFS 5015), NORV (strain 0234), POTV (strain BeAr7272) and TENV (strain A9-171b). CVV-478 was originally isolated from mosquitoes collected in Mexico (Farfan-Ale and others 2009, Blitvich and others 2012). All other viruses were obtained from the World Arbovirus Reference Collection at the University of Texas Medical Branch in Galveston, Texas. PRNTs were performed using all BUN serogroup viruses that occur in the USA (as opposed to only those associated with disease) because antibodies to one BUN serogroup virus often cross-react with other viruses in this serogroup due to their close antigenic relatedness (Hunt and Calisher 1979); thus, the exclusion of one or more viruses may result in serological misdiagnosis. PRNTs were performed in six-well plates containing confluent monolayers of African Green Monkey kidney (Vero) cells following published protocols (Beaty and others 1995). Initially, all sera were screened at a single dilution of 1:20. All sera that tested positive for BUN serogroup-specific antibodies were further diluted and tested by PRNT in order to identify the viruses responsible for these infections. Titres were expressed as the reciprocal of highest serum dilutions yielding $\geq$90 per cent reduction in the number of plaques (PRNT$_{90}$). For aetiological diagnosis, the PRNT$_{90}$ antibody titre to the respective virus was required to be at least four-fold greater than that to the other viruses tested.

Arthropod collections

Arthropods were collected at 10 trap sites in eight Montana counties from May to August 2013 (Fig 1). Five arthropod collection sites were adjacent to or within 5 km of summer-pastured sheep. Collections were made using Centers for Disease Control and Prevention (CDC) light traps placed near the edges of stock ponds or other lentic water sources. Traps were baited with dry ice in the late afternoon, and arthropods were collected early the following morning and transported to the laboratory on dry ice. C. sonorensis were identified according to morphological characteristics and sorted into pools of up to 50 according to date and study site.

Homogenisations

C. sonorensis were placed in polypropylene, round-bottom 5 ml tubes with 1.8 ml phosphate-buffered saline (pH 7.4) supplemented with 100 units/ml penicillin, 100 μg/ml streptomycin and 2.5 μg/ml fungizone. Four 4.5 mm diameter copper-clad steel beads (BB-calibre airgun shot) were added to each tube, and midge pools were homogenised by vortexing for 60 seconds. Homogenates were centrifuged (3000 rpm, 10 minutes, 4°C) and supernatants were collected.

Virus isolation in cell culture

An aliquot (100 μl) of each homogenate was added to 0.5 ml of maintenance medium which consisted of Dulbecco’s modified Eagle medium (DMEM; Invitrogen, Carlsbad, California, USA) supplemented with 2 per cent fetal bovine serum, 2 mM L-glutamine, 100 units/ml penicillin, 100 μg/ml streptomycin and 2.5 μg/ml fungizone. Supernatants were filtered using a 0.22 μm filter and inoculated onto subconfluent monolayers of Vero cells in 6-well plates. Cells were incubated for one hour at room temperature on an orbital shaker and then 5 ml of maintenance medium was added to each well. Cells were incubated at 37°C in the presence of 5 per cent CO$_2$ for 14 days and monitored regularly for cytopathic effect.

Reverse transcription PCR

Total RNA was extracted from homogenates using the QIAamp viral RNA extraction kit (QIAGEN, Valencia, California, USA) and analysed by RT-PCR using primers BCS82C and BCS332V, which are specific to orthobunyaviruses in the BUN and California serogroups (including CVV, MDV, POTV and TENV (Kuno and others 1996) and LOKV and NORV (B.J. Blitvich, unpublished data)). Complementary DNAs were generated using Superscript III reverse transcriptase (Invitrogen), and PCRs were performed using Taq polymerase (Invitrogen).

RESULTS

BUN serogroup virus-specific antibodies were identified in 41 (39.4 per cent) sheep (Table 1). Of these, three were seropositive for MDV, one was seropositive for CVV, one was seropositive for LOKV and 36 had antibodies to an undetermined BUN serogroup virus. Sixteen sheep that had antibodies to an undetermined BUN serogroup virus had low PRNT$_{90}$ titres to MDV and no detectable
antibody titres to any of the other viruses. Antibodies to NORV, POTV and TENV were not identified in any sheep.

A total of 30,606 *C. sonorensis* were collected and sorted into 629 pools. All were homogenised and an aliquot of each homogenate was tested by virus isolation in Vero cells. All homogenates were negative for cytopathic virus. A subset of homogenates (n=200) was further tested by RT-PCR using primers specific to BUN and California serogroup viruses. All homogenates were negative.

**DISCUSSION**

BUN serogroup virus-specific antibodies were identified in 41 (39.4 per cent) sheep. At least three viruses were responsible for these infections: CVV, LOKV and MDV. None of these viruses have been previously reported in Montana. CVV has been detected throughout much of North America including North and South Dakota which border Montana to the east, and Alberta and Saskatchewan which border Montana to the north (Iversen and others 1979, Calisher and others 1986, Pabbaraju and others 2009) but there are no published data demonstrating the occurrence of this virus in Montana. Previously, LOKV activity had only been reported in California, Colorado, New Mexico, Texas and Utah while MDV is known to occur in all of the aforementioned states as well as Arizona (Crane and others 1983, Calisher and others 1986, Kramer and others 1990). Our findings suggest that the geographic distribution of CVV, MDV and LOKV is wider than previously reported.

Because our data indicate that BUN serogroup viruses commonly infect sheep in Montana, it is likely that some of the sheep analysed in this study had been exposed to more than one BUN serogroup virus. However, information on the antibody responses in vertebrates sequentially infected with BUN serogroup viruses is limited and research should be performed to address this issue. There is currently only one report that describes the antibody responses in vertebrates experimentally inoculated with different BUN serogroup viruses. The study was performed using white-tailed deer that were first inoculated with CVV or POTV then sequentially challenged with the alternate virus five to seven months later (Blackmore and Grimstad 1998).

As already noted, 36 sheep had antibodies to an undetermined BUN serogroup virus. Of these, 16 had low PRNT90 titres to MDV and no detectable antibody titres to any of the other viruses. This could indicate that the sheep had been infected with MDV several years ago and, because neutralising antibody levels steadily decline over time (Gibbs and others 2005), only trace amounts of MDV-specific antibodies remained. Alternatively, the sheep could have been infected with a novel BUN serogroup virus that is more closely related to MDV than it is to the other viruses included in the PRNTs. In this regard, the three sheep seropositive for MDV had MDV PRNT90 titres that could be considered low (all titres were 80); thus, we cannot dismiss the possibility that these sheep had instead been infected with an unrecognised MDV-like virus and that the MDV PRNT90 titres are a consequence of serological cross-reactivity. Likewise, the sheep seropositive for LOKV had a LOKV PRNT90 titre of 80 and, therefore, an unknown LOKV-like virus may have been responsible for this infection.

BUN serogroup viruses were not isolated from any *C. sonorensis* collected in this study. Our rationale for testing the midges by virus isolation in cell culture is because this technique is not restricted to the isolation of BUN serogroup viruses; other arthropod-transmitted viruses can also be detected. However, viral RNA is more stable than
infectious viral particles and therefore a subset of *C. sonorensis* was further tested by RT-PCR using BUN serogroup-reactive primers. Another reason why we also decided to perform RT-PCRs is because this technique has the potential to detect novel BUN serogroup viruses that lack the capacity to replicate in Vero cells. However, as already noted, viral RNA was not detected in any midges.

In summary, we provide evidence of BUN serogroup virus infection in sheep but not *C. sonorensis* in Montana. At least three viruses were responsible for these infections: CVV, LOKV and MDV. This study provides the first evidence of CVV, MDV and LOKV activity in Montana. Additional research is needed to determine the impact of these viruses on ovine health.

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**TABLE 1:** PRNT data for sheep with Bunyamwera serogroup virus-specific antibodies

| Serum ID | PRNT<sub>90</sub> titre | CVV | LOKV | MDV | NORV | POTV | TENV | PRNT diagnosis |
|----------|--------------------------|-----|------|-----|------|------|------|----------------|
| A03      | –                        | 40  | –    | –   | –    | –    | –    | *Und          |
| A13      | –                        | 40  | –    | –   | –    | –    | –    | Und           |
| A15      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A17      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A21      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A25      | ≥1280                    | 160 | 40   | 160 | –    | 20   | –    | CVV           |
| A30      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A34      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A35      | –                        | 40  | –    | –   | –    | –    | –    | Und           |
| A36      | –                        | 40  | –    | –   | –    | –    | –    | Und           |
| A37      | –                        | 40  | –    | –   | –    | –    | –    | Und           |
| A39      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A41      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A42      | –                        | 80  | –    | –   | –    | –    | –    | MDV           |
| A44      | –                        | 40  | –    | –   | –    | –    | –    | Und           |
| A45      | –                        | 80  | –    | –   | –    | –    | –    | MDV           |
| A49      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A53      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| A60      | –                        | 40  | –    | –   | –    | –    | –    | Und           |
| B01      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| B05      | –                        | 40  | –    | –   | –    | –    | –    | Und           |
| B07      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| B13      | 640                      | 320 | 160  | 40  | 20   | 160  | –    | Und           |
| B14      | –                        | 80  | –    | –   | –    | 20   | –    | LOKV          |
| B17      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| B18      | –                        | 20  | 20   | 20  | –    | 20   | –    | Und           |
| B19      | –                        | 20  | –    | –   | 20   | –    | –    | Und           |
| B20      | –                        | 80  | 20   | 20  | 20   | –    | –    | MDV           |
| B21      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| B22      | –                        | 20  | –    | –   | 20   | –    | –    | Und           |
| B24      | 40                       | 20  | 20   | 20  | 20   | 20   | 20   | Und           |
| B25      | –                        | 20  | –    | –   | –    | 20   | –    | Und           |
| B29      | –                        | 20  | –    | –   | 20   | –    | –    | Und           |
| B30      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| B31      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| B32      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| B35      | –                        | 20  | 20   | –   | –    | –    | –    | Und           |
| B36      | –                        | 20  | –    | –   | –    | –    | –    | Und           |
| B37      | –                        | 20  | –    | –   | 20   | –    | –    | Und           |
| B38      | –                        | 20  | 20   | 20  | 20   | 20   | –    | Und           |
| B39      | –                        | 20  | 20   | 20  | 20   | –    | –    | Und           |

*<20
†Undetermined Bunyamwera serogroup virus
CVV, Cache Valley virus; LOKV, Lokern virus; MDV, Main Drain virus; NORV, Northway virus; POTV, Potosi virus; PRNT, plaque reduction neutralisation test; TENV, Tensaw virus

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Contributors GDJ and DGH coordinated the collection of sera and midges, CSB and ZNC performed the PRNTs, PI tested midges by virus isolation and RT-PCR, FJP and LCB participated in the design and coordination of the study, and JB conceived of the study, participated in its design and coordination and drafted the manuscript. All authors read and approved the final manuscript.

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