

Schmallenberg virus:

Is there evidence the virus continues to circulate?



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Conclusions Low level of SBV circulation in 2013 and 2014

No evidence SBV circulation in 2015

Introduction

Schmallenberg virus (SBV), transmitted by *Culicoides* spp. biting midges, causes ruminant abortions and congenital malformations, and mild clinical signs in adult dairy cattle (milk drop, fever, diarrhoea).

Following the initial Schmallenberg epidemic (2011 – 2012) the number of newly reported cases rapidly declined suggesting that virus circulation had ceased to occur since 2012. Recently (2014), SBV re-emergence has been reported in Germany¹. However, there is limited research on SBV circulation since 2012.

Objectives

Investigate SBV circulation in dairy herds in 2013, 2014 and 2015

Evidence of SBV circulation

Winter 2014

- 95.5% of animals (95.6% of cows and 99.1% of heifers) re-sampled in winter 2014 remained seronegative after the 2014 vector-active season (Fig. 2)
- 22 animals (0.9%; 10 cows, 5 heifers and 7 youngstock) originating in 17 herds (1-4 animals per herd) tested seropositive

Winter 2015

• All youngstock tested seronegative, including the 7 test-positive animals from winter 2014

Culicoides midge vector spp.

following initial exposure to the virus in 2012

Determine the extent of the susceptible animal population in previously exposed dairy herds

Materials and Methods

✓ Herds

26 Irish dairy herds (exposed to SBV in 2012) Herd size range 58-444 lactating animals per herd

✓ Blood samples

Individual animal blood samples collected before (spring) and following (winter) the 2014 vector-active season and following (winter) the 2015 the vector-active season were tested for SBV-specific antibodies (ID Screen[®] ELISA *Schmallenberg virus Competition Multi-species*)

Blood	Spring 2014	Winter 2014	Winter 2015
samples	n= 5531	n= 2483	n= 1440
Lactating cows	4047	288*	
(born before 2013)	(73%)	(12%)	
Heifers	1480	645*	_
(born spring 2013)	(27%)	(26%)	
Youngstock		1550	1440
(born spring 2014)		(62%)	(100%)

*Blood samples collected from a subsample (n = 933) of seronegative cows and heifers identified in spring 2014

All 138 *Culicoides* pools tested negative for SBV RNA \bullet

Fig. 2 Animal-level seroprevalence after the 2014 vector-active season



✓ *Culicoides* SBV vectors spp.

3,043 pigmented *Culicoides* SBV vector spp. (138 pools; average 22 *Culicoides* specimens per pool) collected from 10 study herds during the 2014 vector-active season were analysed for SBV RNA using rt-RT-PCR

Results

Spring 2014

- Animal-level seroprevalence was 62.5%, 84.7% in cows and 0.6% in heifers
- Within-herd seroprevalence ranged widely from 8.5% to 84.1% in the 26 herds (Fig. 1); 10.7% to 100% in cows and 0.2% to 1.4% in heifers

Fig. 1 Within-herd seroprevalence in Spring 2014



Positive Negative Inconclusive

Discussion

- \succ A wide variation in within-herd seroprevalence in spring 2014 suggests varying risks of infection before the 2014 vector-active season
- \succ The 22 test-positive animals (0.9%; 10 cows, 5 heifers and 7 youngstock) identified in winter 2014 originated in 17 herds (1-4 animals per herd)
- All Culicoides pools tested negative for SBV RNA
- > The 7 test-positive youngstock identified in winter 2014 tested seronegative in winter 2015, suggesting their initial positive result was due to persistence of maternal antibodies
- > These findings suggest SBV recirculated at a low level in these herds in 2013 and 2014, while there was no evidence of SBV circulation in youngstock in 2015. A large population of naive animals, youngstock in particular, were identified in these herds and may be at risk of SBV infection should SBV re-emerge as it has done in continental Europe

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