

Follow-up report 1 (Final report)

16/04/2021

The event is resolved. No more follow-up reports will be submitted.

Sender Country/territory Report ID

Delegate of Spain Spain FUR_149554

Report reference Event status Self-declaration

Cierre Sars-Cov GAL Resolved No

General information

Country or zone - Country Disease - SARS-CoV-2 in animals (Inf. Started on - 03/12/2020

with)

Animal type - Terrestrial Confirmed on - 19/01/2021 Causal agent - SARS-CoV-2

Ended on - 14/02/2021 Disease category - Emerging Reported on - 16/04/2021

Reason - Emerging disease

Disease impact

Outbreak morbidity - 0.32 Outbreak mortality - 0.0

Zoonotic potential - Yes Zoonotic potential description - The infected animals showed no

clinical sign

Epidemiology

Source of the event or origin of the infection - Unknown or inconclusive

Epidemiological commentsLikely introduction through contact of animals with infected workers

Control measures at event level

Domestic control measures

Applied

- Traceability

- Surveillance within infected and/or protection zone
- Surveillance within infected and/or protection zone
- Surveillance within containment and or the protection zone
- Surveillance outside infected and/or protection zone
- Surveillance outside infected and/or protection zone
- Surveillance outside containment and or the protection zone
- Screening
- Process to inactivate the pathogenic agent in products or byproducts
- Official disposal of carcasses, by-products and waste

To be applied

- Stamping out
- Disinfection / Disinfestation

Wild control measures

Applied

- Traceability
- Surveillance within infected and/or protection zone
- Surveillance within infected and/or protection zone
- Surveillance within containment and or the protection zone
- Surveillance outside infected and/or protection zone
- Surveillance outside infected and/or protection zone
- Surveillance outside containment and or the protection zone
- Screening
- Official disposal of carcasses, by-products and waste

To be applied

- Stamping out
- Disinfection / Disinfestation

- Process to inactivate the pathogenic agent in products or byproducts

Diagnostic

Clinical signs - No

Method of diagnostic - Diagnostic test

Test name	Category	Test type	Laboratory	Species sampled	Outbreaks	Tested from	Tested until	Result
Real-time polymerase chain reaction (real-time PCR)	Nucleic acid detection	Laboratory Test	Instituto de Investigaciones Marinas de Vigo – Consejo Superior de Investigaciones Científicas (CSIC), ESP	Mustelidae		19/01/2021		Positive
Real-time polymerase chain reaction (real-time PCR)	Nucleic acid detection	Laboratory Test	Laboratorio Central de Veterinaria de Algete	Mustelidae		19/01/2021		Positive

Quantitative data summary

Measuring unit - Animal

Species	Туре	Susceptible	Cases	Deaths	Killed and disposed of	Slaughtered	Vaccinated	Outbreak morbidity	Outbreak mortality
Mustelidae	New	-	-	-	3100	-	0	-	-
Mustelidae	Total	3100	10	0	3100	0	0	0.32	0.0
All species	New	-	-	-	3100	-	0	-	-
All species	Total	3100	10	0	3100	0	0	0.32	0.0

Event morbidity - 0.32 Event mortality - 0.0

Morbidity range - 0.32-0.32 Mortality range - 0-0

Outbreaks

1000143912-A Baña

Outbreak reference - 2021/1 Started on - 03/12/2020 First administrative division - Galicia

Ended on - 14/02/2021 Second administrative division - A Coruña Epidemiological unit - Farm

Geographic coordinates - Third administrative division - n.a. (145) Location - A Baña

42.9834,-8.7851

Description of the affected populationFarm with a census of 3,100 adult American mink, of which 2,500 are females and 600 males.

Species	Туре	Measuring unit	Susceptible	Cases	Deaths	Killed and disposed of	Slaughtered	Vaccinated
Mustelidae	New	Animal	-	-	-	3100	-	0
Mustelidae	Total	Animal	3100	10	0	3100	0	0
All species	New	Animal	-	-	-	3100	-	0
All species	Total	Animal	3100	10	0	3100	0	0

Map legend **OUTBREAKS** REPRESENTATION ON MAP

OUTBREAKS NATURE

Domestic species

Wild species

Domestic & Wild

OUTBREAKS STATUS

Outbreak currently reported

Ongoing outbreak

Resolved outbreak

AGGREGATION REPRESENTATION

< 20 outbreaks

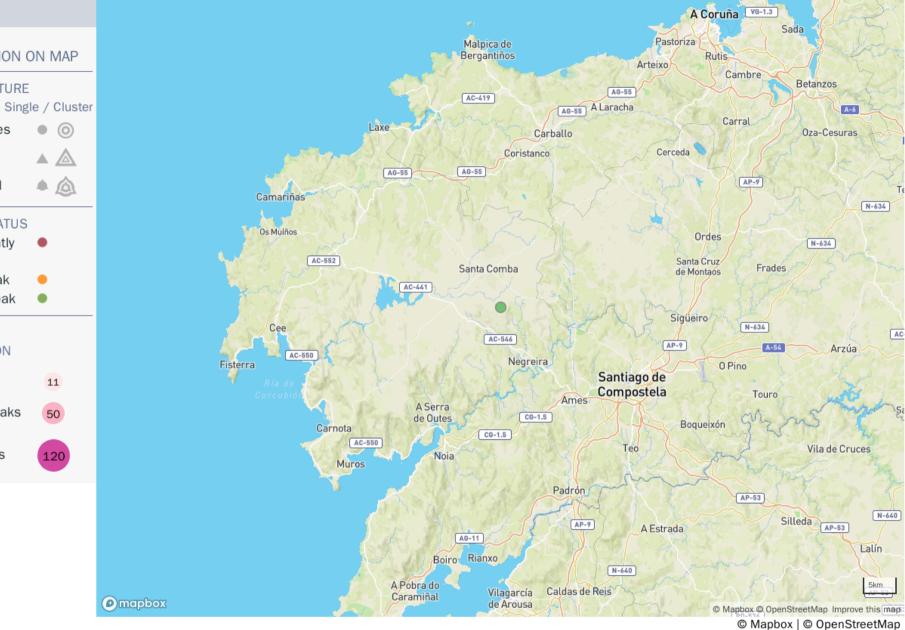
20 - 100 outbreaks

> 100 outbreaks

120

11

50



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