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Exposure of harbour seals *Phoca vitulina* to *Brucella* in declining populations across Scotland

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ABSTRACT: Since 2000 there has been a major decline in the abundance of Scottish harbour seals *Phoca vitulina*. The causes of the decline remain uncertain. The aim of this study was to establish the extent to which the seals in the regions of greatest decline have been exposed to *Brucella*, a bacterial pathogen that causes reproductive failure in terrestrial mammalian hosts. Tissues from dead seals collected between 1992 and 2013 were cultured for *Brucella* (n = 150). Serum samples collected from live capture–released seals (n = 343) between 1997 and 2012 were tested for *Brucella* antibodies using the Rose Bengal plate agglutination test (RBT) and a competitive enzyme-linked immunosorbent assay (cELISA). In total, 16% of seals cultured had *Brucella* isolated from one or more tissues, but there were no pathological signs of infection. The cELISA results were more sensitive than the RBT results, showing that overall 25.4% of seals were seropositive, with the highest seroprevalence in juveniles. As there was no evidence of either a higher seroprevalence or higher circulating antibody levels in seropositive animals in the areas with the greatest declines, it was concluded that *Brucella* infection is likely not a major contributing factor to recent declines. However, the consistently high proportion of seals exposed to *Brucella* indicates possible endemicity in these populations, likely due to *B. pinnipedialis*, which has demonstrated a preference for pinniped hosts. Importantly, given the close proximity between seals, humans and livestock in many areas, there is the potential for cross-species infections.

KEY WORDS: Pinnipeds · *Brucella* · Disease · Cultures · Seroprevalence · Antibodies · ELISA · Rose Bengal plate agglutination test

INTRODUCTION

Aerial surveys have been carried out by the Sea Mammal Research Unit, to monitor harbour seal *Phoca vitulina* populations around Scotland since 1985, and declines in a number of these populations have been seen since 2000 (Lonergan et al. 2007). Major declines of 68% in Orkney, 50% in Shetland, and 90% in the Firth of Tay have been documented in particular (SCOS 2012). However, the pattern of the declines is not universal, as some areas remain more stable while the populations in other areas con-

tinue to decrease in size. The population in the Eden and Firth of Tay Special Area of Conservation, for example, has experienced the most dramatic and sustained declines of over 90% in the last 15 yr, with a most recent estimate of just 29 individuals left in 2014 (Hanson et al. 2017).

Many potential causes of the decline have been suggested, but the contributing factors remain uncertain. Some of these include predation by killer whales (Bolt et al. 2009), competition for food with other marine top predators (SCOS 2012), exposure to biotoxins from harmful algal blooms (Hall & Frame

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2010), deliberate shooting (Thompson et al. 2007), accidental mortalities as a result of interactions with shipping vessels (Thompson et al. 2010) and predation by grey seals *Halichoerus grypus* (Brownlow et al. 2016). A further potential contributing factor to these declines is infectious disease, but there has been a lack of reports of sick animals by the Scottish Marine Animals Strandings Scheme (SMASS) or the Scottish Society for the Prevention of Cruelty to Animals (SSPCA). This suggests that if infectious disease was present in these populations, and was contributing to the observed declines, it could either be affecting the reproductive success of the animals, or causing them to die very quickly once infected, or both. A particular infectious agent of interest in this respect is *Brucella* as it is known to cause reproductive failure in other mammalian hosts. The aim of this study was to establish the extent to which harbour seals in Scotland have been exposed to *Brucella* over time, both before and during the observed population declines, and whether this could be a potential contributing factor to the major declines in some areas.

Members of the genus *Brucella* are Gram-negative, rod-shaped bacteria that cause chronic disease most commonly associated with abortions and infertility in domestic livestock (Seleem et al. 2010). Since the first reports of *Brucella* in a marine mammal in 1994 (Ross et al. 1994), infections have been recognised in a range of pinniped and cetacean species worldwide (Thakur et al. 2012). Strains isolated from marine mammals have been shown to be phenotypically and genetically distinct from those isolated from terrestrial mammals, and 2 species have been described that have pinnipeds and cetaceans as their preferred hosts: *B. pinnipedialis* and *B. ceti*, respectively (Foster et al. 2007). *B. ceti* infections in cetaceans have been associated with various pathologies which include abortions and neonatal mortality (Miller et al. 1999), epididymitis in males (Dagleish et al. 2008), meningoencephalitis (González et al. 2002, Jauniaux et al. 2010, Alba et al. 2013, Garofolo et al. 2014), abscesses (Foster et al. 1996, Foster et al. 2002), endocarditis (González-Barrientos et al. 2010), mastitis, pneumonia, peritonitis, osteomyelitis and spinal discospondylitis (Foster et al. 2002). In contrast, pathology associated with *B. pinnipedialis* in seals is lacking despite several reports of its isolation (Foster et al. 2002, Tryland et al. 2005, Nymo et al. 2011, Siebert et al. 2017). With respect to harbour seals specifically, *Brucella* has previously been isolated from wild animals (Ross et al. 1994, Garner et al. 1997, Foster et al. 2002, Prenger-Berninghoff et al. 2008), and in several areas they have also been found

to be seropositive (Ross et al. 1996, Maratea et al. 2003, Gaydos et al. 2005, Hueffer et al. 2013). However, the extent to which harbour seals in Scotland, or the United Kingdom in general, have been exposed to this pathogen since the onset of the declines is unknown. Here, tissue samples from dead harbour seals collected by the SMASS over 20 yr were cultured to test for *Brucella* isolates. In addition, harbour seal serum samples from live capture–release studies by the Sea Mammal Research Unit over a 14 yr period were tested for *Brucella* antibodies. Tempo–spatial patterns in seroprevalence were examined in order to investigate the potential role of *Brucella* as a contributing factor to the Scottish harbour seal declines.

MATERIALS AND METHODS

Brucella cultures

Tissue samples from 150 dead harbour seals from across Scotland which had received post mortem examination were collected by the SMASS between 1992 and 2013 as part of systematic surveillance studies. Microbiological culture was performed, including specific methods for *Brucella* isolation. The selected tissues varied between animals but typically included lung, liver, kidney, spleen and small intestine, but also brain, pancreas, reproductive tissue, various lymph nodes and any abscesses apparent at post mortem. These were collected from approximately equal numbers of males (n = 61), females (n = 41) and unsexed animals (n = 48). The majority of cases were adult animals although some juveniles were also sampled. Tissues were processed using a standardised method (Foster et al. 2002) and cultured on Columbia sheep blood agar (CSBA; Oxoid) and Farrell's medium incubated at 37°C in air with 5% added CO₂. Isolates with colonial appearance typical of *Brucella* on either medium were identified as *Brucella* using phenotypic tests as previously described (Foster et al. 2002). A multi-locus variable number of tandem repeats analysis (MLVA-16) was used to confirm species designation (Maquart et al. 2009).

Serum sampling procedure

Blood samples were collected from 343 live-captured harbour seal adults, juveniles (<50 kg and/or 120 cm) and suckling pups from multiple haul out sites around 5 areas of Scotland: the South East (the

Eden Estuary, the Firth of Tay and the Firth of Forth), the North East (the Moray Firth, Dornoch Firth, Loch Fleet and the Pentland Firth), the North West (the Isle of Skye, Loch Shieldaig, and the Loch Nan Uamh Islands), the South West (the Sound of Jura and south east Islay), and Orkney (Fig. 1). Samples were collected between 1997 and 2012 at varying times of the year. Due to the opportunistic nature of the analysis of stored samples, sample sizes varied regionally, across years and between age classes, although the male to female ratio was approximately equal with 182 males and 160 females (Table 1). The seals were captured in nets on haul outs or in the water, and were sedated with Zoletil 100 (Virbac) at a dose rate of 0.5 ml per 100 kg body weight intravenously. Blood samples were taken from the extradural vein immediately after the immobilisation of the animal. The whole blood samples were spun, sera was collected, and aliquots were frozen at -20°C for later analysis. Samples were collected under the Animal (Scientific Procedures) Act, 1986, Home Office Pro-

ject and Personal Licences issued to the Sea Mammal Research Unit.

Serological methods

In a preliminary trial, the Rose Bengal plate agglutination test (RBT) was used to test stored serum samples for the presence of *Brucella* antibodies. Serum samples were tested with the Micropath Rose Bengal kit against the *B. abortus* antigen (Omega Diagnostics) following the kit instructions. Samples were either classed as positive or negative based on visually discernible agglutination of antigens. A positive and a negative control supplied by the kit were used for each set of 4 serum samples tested simultaneously.

Following the completion of the Rose Bengal trials with the successful detection of antibodies, the variation in seroprevalence, i.e. the proportion of animals with antibody levels higher than a background

threshold level set for terrestrial mammals, as well as the absolute antibody levels in the samples, were investigated using a competitive ELISA (cELISA). Here, polystyrene microtitre plates were coated with *Brucella melitensis* lipopolysaccharide (LPS) antigen. Both positive (positive serum from an experimentally infected goat with *B. melitensis*) and negative controls (no sera) were tested. Serum and a peroxidase-labelled monoclonal antibody (BM40 from a locally held hybridoma) were added to the plates and incubated for 30 min at room temperature. The plate was then washed, and chromogen and substrate were added and incubated for a further 15 min at room temperature, shaking at 160 rpm. The plates were read at an optical density (OD) of 450 nm. The mean OD of duplicate wells was expressed as a percentage of antibodies binding to the plate. Test samples with an OD of less than 60% of the conjugate-only control (no sera added) were recorded as positive. This cut-off threshold was established based on serology results from terrestrial mammals (Perrett et al. 2010). As such, a weak reaction, indicative of low antibody levels, was considered to be between 30 and 60% antibody binding, while a strong reac-

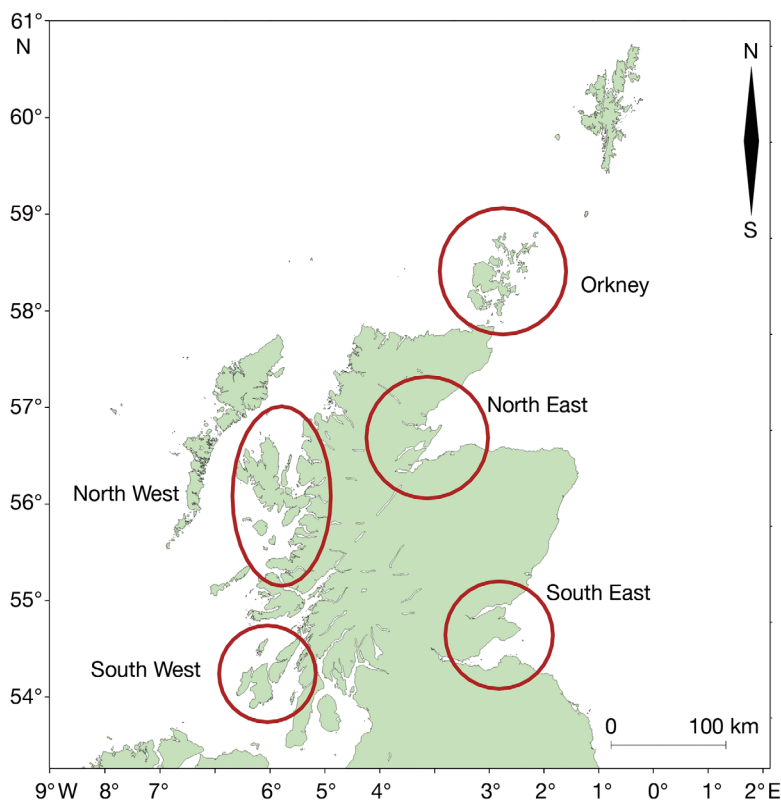


Fig. 1. Sampling regions of live-captured harbour seals across Scotland between 1997 and 2012 by the Sea Mammal Research Unit. Serum samples were grouped into 5 main regions across Scotland as indicated by the circles on the map. Over this time period, the populations along the west coast, marked as 'North West' and 'South West' were stable while the populations in 'Orkney', the 'North East' and the 'South East' underwent precipitous declines (SCOS 2012)

Table 1. Serological samples collected from harbour seals across the 5 sampling regions of Scotland and over 12 yr broken down into 4 time periods. Samples are grouped by sex and age class. A total of 306 adults, 15 juveniles and 22 pups were sampled

Sampling period	Sex	Age class	Serological samples collected by region				
			North East	North West	Orkney	South East	South West
1997–2000	Male	Adults	–	–	–	37	–
		Juveniles	–	–	–	1	–
		Pups	–	–	–	3	–
	Female	Adults	–	–	–	21	–
		Juveniles	–	–	–	1	–
		Pups	–	–	–	–	–
2001–2005	Male	Adults	5	–	11	15	14
		Juveniles	–	–	–	1	–
		Pups	–	–	–	–	–
	Female	Adults	10	–	12	9	4
		Juveniles	1	–	–	2	2
		Pups	–	–	–	–	1
2006–2008	Male	Adults	3	1	10	16	8
		Juveniles	–	–	–	–	1
		Pups	–	–	–	–	5
	Female	Adults	7	1	15	6	7
		Juveniles	1	–	–	–	–
		Pups	2	–	11	–	–
2009–2012	Male	Adults	1	14	21	10	2
		Juveniles	1	–	1	–	1
		Pups	–	–	–	–	–
	Female	Adults	6	11	20	1	8
		Juveniles	1	–	–	–	1
		Pups	–	–	–	–	–

tion, indicative of high circulating antibody levels, was considered to be <30 %.

Statistical analysis of serological data

All statistical analyses were performed using the statistical package R, version 3.1.2 (R Core Development Team 2014). Results were considered statistically significant at $p \leq 0.05$. Two different statistical approaches were taken to investigate firstly, variation in the seroprevalence data, and secondly, variation in the antibody levels in the seropositive individuals.

For the seroprevalence data, generalised linear models (GLMs) with a binomial distribution were fitted to the seroprevalence data with individuals classed as seropositive (1) and seronegative (0) for the 2 tests separately. The 14 yr of data were split into 4 time periods: 1997 to 2000 ($n = 63$) represents the years before the start of the decline, and the years be-

tween 2001 and 2012 were split into 3 periods with approximately equal numbers of samples in each to give the maximum statistical power for the analysis. These were 2001 to 2005 ($n = 87$), 2006 to 2008 ($n = 94$) and 2009 to 2012 ($n = 99$). A global model including all explanatory variables of interest (region, sex, age class, time period and an interaction between region and time period) was generated, and backwards variable selection using the 'step' function in the 'car' library in R v.2.11.1 was performed to identify the combination of variables that best explained the variation in the data by producing the model with the lowest Akaike's information criterion (AIC) value.

In addition, variation in the levels of circulating *Brucella* antibodies were investigated in the seropositive individuals identified using the cELISA data. Antibody binding results of only the seropositive seals were modelled using a GLM with a gamma distribution and a log-link function to model the non-normal distribution of the antibody binding data, as most individuals had low circulating antibody levels while few were very high. Again, a global model with region, sex, age class, time and an interaction

between region and time was generated and backwards variable selection using the 'step' function was used to identify the combination of variables that best explained the variation in the data.

RESULTS

Brucella cultures

Of the 150 animals examined bacteriologically between 1992 and 2013, *Brucella* was isolated from the tissues of 24 individuals (16 %). Details for 11 of these animals have been reported previously (Foster et al. 2002). None of the culture-positive animals showed any signs of pathological lesions associated with infection, and the cause of death was always associated with starvation, trauma or some other viral or bacterial infection, but not *Brucella*. Of the tissues cultured, the 2 that were the most commonly culture-positive were lung (45.8 %) and spleen (41.7 %),

Table 2. Details of 24 *Brucella* culture-positive harbour seals. The last 2 digits of the reference number indicate the year of stranding. (*) Indicates individuals that were culture-positive but seronegative. MLN: mesenteric lymph node; IILN: internal iliac lymph node; EILN: external iliac lymph node; GLN: gastric lymph node; ManLN: mandibular lymph node; HLN: hepatic lymph node; TLN: thoracic lymph node; CLRN: colorectal lymph node; SI: small intestine

Reference no.	Positive cultures	Negative cultures
M2357/93	Spleen	Lung, liver
M2466/93	Spleen	Lung, MLN
M2533/93	Spleen	MLN, SI
M292/94 *	Spleen	Testes, MLN, SI
M336/94 *	IILN	Spleen, MLN
M339/94	GLN	Spleen, IILN
M972/94	EILN, manLN	Spleen, MLN
M490/95	EILN, HLN, IILN, TLN	Lung, spleen, brain, CRLN, GLN, manLN, blood, SI
M514/96	Lung	
M445/99	Lung	Liver, spleen, kidney, brain, MLN, blood, SI
M13/01	Lung	Liver, spleen, kidney, blood
M250/02	Lung, liver, spleen, kidney, MLN, blood	Brain, SI
M305/02	Spleen	Lung, liver, kidney, brain, MLN, cellulitis, SI
M342/02	MLN	Lung, liver, spleen, kidney, brain, blood
M374/02	Lung, liver, spleen, kidney	Brain, MLN
M449/02	Lung	
M599/02	MLN	Lung, liver, spleen, kidney
M43/09	Lung, liver, spleen, kidney, MLN, SI	
M91/10	Lung, brain	Liver, spleen, kidney, MLN
M228/10	Pancreas	Lung, liver, spleen, kidney, brain, MLN, SI
M244/10	Lung, liver, spleen, brain, MLN, SI	Kidney
M273/10	Lung, MLN	Liver, spleen, kidney, brain, pre-scapular LN, SI
M341/11	Lung, spleen, MLN, SI	Liver, kidney, brain, abscess
M337/13	Kidney, brain	Lung, liver, spleen

although not all tissues were sampled consistently across individuals (Table 2). MLVA-16 analysis identified isolates as *B. pinnipedialis* belonging to 1 of 2 sequence types: ST 24 or ST 25. Serum was sampled from 12 of these culture-positive seals, 2 of which were seronegative and were sampled from healthy animals that had been shot (Table 2).

Serology

Test performance

The RBT trials were able to detect antibodies in the archived serum samples, and results showed that across all study sites over the whole sampling period, the prevalence of *Brucella* antibodies was 15.9%. However, the cELISA results showed a higher overall seroprevalence of 25.4%. The prevalence across all age and sex classes, as well as across regions and over time was lower for the RBT results compared to the cELISA results (Table 3).

All of the samples that were positive using the RBT were also classed as positive using the cELISA. These samples had the lowest antibody binding indi-

Table 3. Comparison of the prevalence (% of seropositive harbor seals) using the Rose Bengal plate agglutination test (RBT) and the competitive ELISA (cELISA). The ELISA results indicate a higher overall prevalence of *Brucella* antibodies in harbour seals than the RBT results

Variable	n	% of seropositive animals	
		RBT	cELISA
Scotland	343	15.9	25.4
North East	38	28.6	28.9
North West	27	14.8	33.3
Orkney	101	12.9	28.7
South East	123	19.8	20.3
South West	54	3.9	24.1
Males	183	14.5	25.7
Females	160	17.4	24.4
Adults	306	14.5	25.2
Juveniles	15	60.0	53.3
Pups	22	4.5	9.09
1997–2000	63	12.3	22.2
2001–2005	87	24.7	26.4
2006–2008	94	12.0	18.1
2009–2012	99	13.7	33.3

cating the highest circulating *Brucella* antibodies. Specifically, the mean antibody binding of the samples classed as positive by the RBT was $45.0 \pm 0.06\%$,

while those classed as negative had a mean antibody binding of $74.0 \pm 0.02\%$ (2-sample *t*-test; $t = 8.79$, $df = 51.54$, $p < 0.0001$). Therefore, it seems that the RBT is only able to detect high antibody levels in the serum samples and is the least sensitive of the 2 serological methods tested. As a result, the RBT results may have underestimated the prevalence of *Brucella* antibodies in these harbour seals (Table 3).

Seroprevalence

Using the cELISA seroprevalence data where individuals were classed as either seropositive or seronegative, the best binomial GLM after backwards variable selection included only age class as an important explanatory variable. Juveniles had a significantly higher seroprevalence than adults and pups (p -values < 0.025), while adults and pups were not significantly different from each other ($p = 0.11$). There were no significant changes in prevalence over time or between regions, and there were equal numbers of seropositive males and females.

Antibody levels

Variation in the levels of circulating *Brucella* antibodies were investigated in the seropositive individuals. Backwards variable selection of the GLM using the cELISA antibody binding results of only the

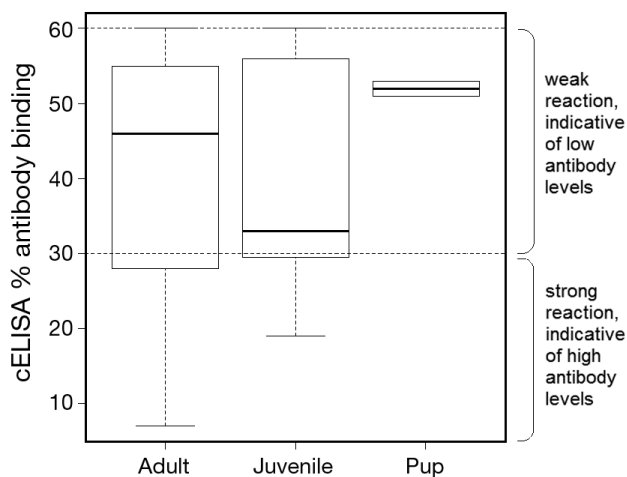


Fig. 2. Competitive ELISA (cELISA) antibody binding of the seropositive harbour seals by age class. Lines in the boxes represent the median values; box edges are the 25th and 75th percentiles. Low antibody binding indicates high levels of circulating *Brucella* antibodies in the seals. Pups had lower circulating antibody levels than both adults and juveniles

seropositive seals revealed that age class and an interaction between region and time period were retained in the final model with significant effects. Pups had near-significant higher antibody binding than both adults and juveniles (both p -values < 0.07), indicating the lowest circulating antibodies in these seropositive individuals (Fig. 2). There was no difference between the circulating antibody levels in juveniles and adults (Fig. 2). The interaction between region and time revealed that there were different patterns in circulating antibody levels in the seropositive seals between regions over the 14 yr sampling period. The highest average circulating *Brucella* antibody levels (shown as the lowest % antibody binding in Fig. 3) were measured in the 2001 to 2005 time period in the South West, and these then decreased over the following years ($p = 0.035$). All areas showed a decrease followed by an increase again over the whole time frame, with the exception of Orkney, which showed a sustained decrease in circulating antibodies between 2001 and 2012 (Fig. 3), and the North West, where seropositive individuals were only recorded in the final time period, but this is likely a reflection of very limited sampling before 2009 (Table 1) rather than a recent introduction of the bacteria to the area. Males and females had similar antibody levels.

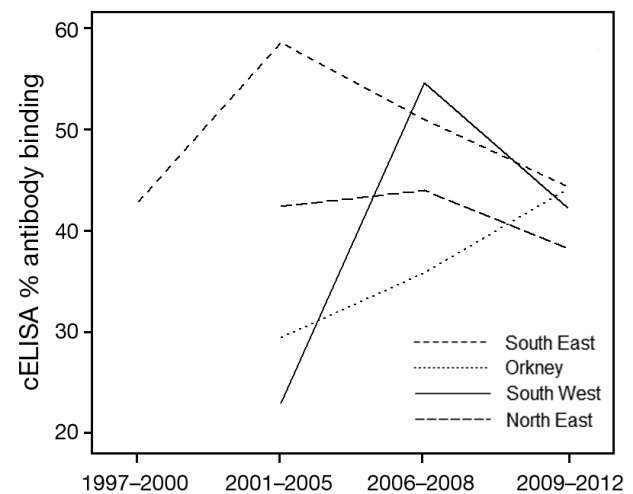


Fig. 3. Interaction plot of the competitive ELISA (cELISA) % antibody binding over time for each sampling region showing variation in seropositive harbour seal individuals. Low antibody binding indicates high levels of circulating *Brucella* antibodies in the seals. The North West sampling region is not included here as seropositive animals were only identified in this area between 2009 and 2012. With the exception of Orkney, the other sampling regions showed a decrease in the circulating levels of *Brucella* antibodies, followed by an increased again over this time period

DISCUSSION

Brucella species were isolated from 16% of the dead stranded animals tested, but there were no signs of *Brucella*-specific pathological lesions associated with infection in these 24 animals. They all appear to have died of other causes, although it is possible that *Brucella* acted as a secondary infection in these cases. The results presented here therefore suggest that harbour seals can be infected by *Brucella*, likely *B. pinnipedialis*, without evidence of associated disease. *B. pinnipedialis* has also been cultured from apparently healthy tissues of a number of other pinniped species, including grey seals *Halichoerus grypus* (Foster et al. 2002, Prenger-Berninghoff et al. 2008), hooded seals *Cystophora cristata* (Foster et al. 2002, Tryland et al. 2005), ringed seals *Pusa hispida* (Forbes et al. 2000) and harp seals *Pagophilus groenlandicus* (Forbes et al. 2000). At present, there is only limited evidence of *Brucella* infection causing disease in any species of phocid seal (Jauniaux et al. 2013), although *Brucella* isolation was suggested as a possible cause of abortion in an otariid species, the California sea lion *Zalophus californianus*, with recovery of *Brucella* from the placenta and stomach contents of an aborted foetus (Goldstein et al. 2009). Extensive typing of a large number of marine mammal *Brucella* strains in a recent study, however, found the Californian sea lion isolates to be similar to *B. ceti* recovered from bottlenose dolphins in the USA and that *B. pinnipedialis* isolates from harbour seals in the USA were found to be closely related to Scottish strains (A. M. Whatmore et al. unpubl. data).

Animal experimentation in cattle with a *Brucella* isolate recovered from a Pacific harbour seal *Phoca vitulina richardsii* resulted in seroconversion and abortion in 2 of 3 pregnant animals, suggesting that some strains of *B. pinnipedialis* may have abortifacient potential (Rhyan et al. 2009). However, there has been no evidence in pinnipeds of disease due to *B. pinnipedialis* as has been seen with *B. ceti* infection in dolphins and porpoises where chronic disease with significant clinical and pathological signs including male infertility, neurobrucellosis, cardiopathies, bone and skin lesions and live strandings have been documented (Guzmán-Verri et al. 2012). While no evidence of disease has been reported in pinnipeds, detecting abortions in wild populations is very difficult, especially if the occurrence remains constant over time and there is limited data on pupping success, as is the case for the populations sampled here. The ability to detect abortions and monitor

pupping success in different populations is therefore needed in order to determine that *B. pinnipedialis* does not cause disease in these seals.

Similar to this study, *Brucella* antibodies have been detected in sera from a number of marine mammal species using Rose Bengal tests (Tryland et al. 1999, 2005, Retamal et al. 2000, Hernández-Mora et al. 2008, Jensen et al. 2013), and using both indirect ELISA (iELISA) and cELISAs primarily designed for ruminants (Tryland et al. 1999, Nielsen et al. 2001, Van Bresseem et al. 2001, Tachibana et al. 2006, Roe et al. 2010, Lynch et al. 2011, Jensen et al. 2013, Nymo et al. 2013a). Here, differences in the prevalence estimates obtained from the RBT and the cELISA results highlight the need to consider test performance when conducting serological studies. It seems that the RBT is only able to detect antibodies when at higher levels, and as such, seropositive samples with low levels of antibody are not recognised.

Our results suggest that the cELISA appears to be a more sensitive test than RBT and is thus able to distinguish between seronegative samples and samples with low antibody levels. The cELISA results were therefore chosen for further analysis over the RBT results as this is thought to be a more robust and objective test. However, when detecting *Brucella* antibodies using serological methods, serological cross-reactions and false positives are potentially a major problem, and may contribute to the higher seroprevalence seen in the cELISA results. It is thought that in cattle most problems caused by cross-reactivity are the result of antibodies produced through the immune response of the animal to other microorganisms sharing similar structural characteristics with the O-polysaccharide of *Brucella* species (Corbel 1985). We cannot rule out the possibility that other cross-reacting bacteria could affect these results. In cattle, it is thought that the cELISA is a more appropriate serological test than the iELISA as it is better able to distinguish between antibodies to *Brucella* species and antibodies to other cross-reacting Gram-negative bacteria (Nielsen 1990, Samartino et al. 1999). In 2 studies on Australian fur seals (Lynch et al. 2011) and Hawaiian monk seals *Monachus schauinslandi* (Nielsen et al. 2005), it was concluded that the iELISA was an unreliable test for the identification of seropositive individuals. Thus, based on this previous evidence, the cELISA was chosen here as an appropriate assay as it is more conservative than an iELISA with a reduced chance of false positives.

The cELISA results indicate that approximately 25% of the seals sampled had antibodies to *Brucella*.

This is within the range of previous studies on harbour seal populations in the North Atlantic, where prevalence ranged between 3.1% (n = 96) in the St Lawrence Estuary, 14% (n = 21) (Maratea et al. 2003) and 50% (n = 8) off the Atlantic coast of the United States (Nielsen et al. 2001). Serology testing of 300 Scottish harbour seals prior to 2002 found 147 (49%) to be positive (Foster et al. 2002). As there appears to have been no change in antibody prevalence over this 14 yr sampling period, or between regions, these data suggest that *Brucella* may be endemic in Scottish harbour seals, and exposure to the bacteria seems to have remained constant over the study period. For endemicity to occur, a high and regular rate of transmission of the bacteria is required within a population.

The transmission of *Brucella* in marine mammals is poorly understood as there is little evidence to support any particular route of infection in these species. It is likely that the routes of transmission are similar to those of terrestrial mammals, whereby transmission occurs through exposure to infected placenta, birth fluids and vaginal secretions as well as by venereal spread (Young 2006). As *Brucella* has been isolated from the reproductive organs of several cetacean species (Miller et al. 1999, Foster et al. 2002, González-Barrientos et al. 2010), and from an aborted foetus of a captive bottlenose dolphin (Ewalt et al. 1994), the most likely mode of transmission of *B. ceti* appears to be through sexual intercourse, vertical transmission from mother to foetus, maternal feeding and contact with aborted foetuses and placental tissues (Guzmán-Verri et al. 2012). The transmission between pinnipeds is even less well understood, but it could be similar to cetaceans. However, transmission may also occur through contact with infected individuals in gregarious species that haul out together in large groups. *Brucella* was cultured from the faeces of a seropositive juvenile harbour seal in captivity (Gaydos et al. 2005), suggesting that some *Brucella*-positive seals are actively shedding the bacteria. In addition, *B. pinnipedialis* was cultured or detected by PCR in harbour seal salivary gland secretions, lungs, urinary bladder and faeces (Lambourn et al. 2013), suggesting that seals could be exposed to the bacterium via exposure to oral secretions, urine or faeces on haul-outs. *Brucella* has also been isolated from subcutaneous lesions in cetaceans (Foster et al. 1996, 2002), so the potential for direct contact with similarly infected skin lesions, should they occur, in pinnipeds that haul out together may present another mode of transfer of the bacteria, although such lesions have not been reported to date.

Together, this could make harbour seals more at risk of bacterial transfer at particular times during their life cycle when they haul-out in larger numbers during the breeding season and during the moult. Thus, the requirements of a high and regular rate of transmission of the bacteria for endemicity to occur could be met for harbour seals.

There was no regional variation in seroprevalence across Scotland, indicating that seals in the declining populations in Orkney and along the East coast have similar proportions of seropositive seals to the stable populations along the West coast. In addition, there has been no change in prevalence over time, even in declining populations, and none of the seals sampled in this study showed any overt signs of ill health. These results further support the hypothesis that they may be infected by a strain of the bacteria that appears to be having little effect on their health. Other pinniped species have also been shown to be seropositive and yet remain apparently healthy and asymptomatic (Nielsen et al. 1996, 2005, Retamal et al. 2000, Nymo et al. 2011). Together, these results indicate that the bacteria may only cause a mild and transient infection, and *B. pinnipedialis* is most likely not a major cause of the harbour seal decline in Scotland. Other potential causes of the declines should therefore continue to be investigated.

Juveniles showed the highest overall prevalence of the 3 age classes. It has been reported that the higher incidence in juveniles may be a result of recent exposure to the pathogen due to a change to a prey-based diet after they are weaned (Lynch et al. 2011, Nymo et al. 2013b). Lungworms carrying *Brucella* in fish prey species may be a means by which marine mammals become infected with the bacterium, as was suggested when *Brucella* was isolated from the lungworms in a harbour porpoise *Phocoena phocoena* (Dawson et al. 2008). *Brucella* species have also been found by immunohistochemical staining in the uterus and the intestinal lumen of female *Parafilaroides* lungworms from a Pacific harbour seal (Garner et al. 1997), and it was postulated that, based on the life cycle of the parasite, the larvae migrate through the respiratory tract and are then swallowed. From there, they pass through the digestive tract and out into the environment in the faeces, where they are taken up by fish and ultimately by the seal. The parasitic larvae are released into the gastrointestinal tract of the animal, and when they mature into adults, they migrate to the lungs and continue the life cycle (Howard et al. 1983). It may also be significant that lung was the body tissue with the highest isolation rate from the 24 harbour seals that were positive by *Brucella* culture.

As such, the high levels of antibodies in seropositive juveniles may suggest a more recent exposure to *Brucella* as they first start to eat fish containing the infected parasites, but it is not necessarily indicative of an active infection. The seropositive adults have high antibody levels which may be indicative of both previous and regular exposure to the bacteria.

The finding that there were lower levels of *Brucella* antibodies in the seropositive pups compared to both seropositive adults and juveniles is surprising. It would be expected that passively transferred maternal antibodies would be present in pups, as they are found in the offspring of antibody-positive mothers in terrestrial species (Ray et al. 1988, Thakur et al. 2002, Rhyan et al. 2009). However, a lower seroprevalence was seen in Australian fur seal (Lynch et al. 2011), Hawaiian monk seal (Aguirre et al. 2007), hooded seal (Nymo et al. 2013b) and Alaskan harbour seal pups (Zarnke et al. 2006) compared to both adults and juveniles. In these studies, it was concluded that pups may have had maternal antibodies at titres lower than the threshold of detection used in their serological tests. These data support the theory that pups likely have low levels of maternal antibodies and that they may not be exposed to infection until a later stage post-weaning (Lynch et al. 2011). These findings further highlight the need for investigations into the timing of first exposure to *Brucella* and seroconversion as well as the development of specific thresholds of detection for antibodies to marine mammal strains of *Brucella* in various serological tests.

While the overall proportion of positive seals did not change across the different sampling regions over time, there were varying patterns of high and low antibody levels measured in the seropositive seals. Higher antibody levels were not recorded in the declining populations, however, and there were no populations with consistently higher or lower antibody levels. The presence of antibodies does not necessarily suggest that the animals had a current or active infection at the time of sampling. The variation over time seen here between populations likely reflects cycles of infection followed by clearance in infected individuals that do not show any clinical signs of the disease. While the apparently high exposure rates of Scottish harbour seals to *Brucella* appear not to be having a negative impact on their populations, such levels may have important implications for cross-species infections between humans and domestic livestock, where infections may lead to disease. Currently, a total of 53 marine mammal species worldwide have been shown to be seropositive for *Brucella* antibodies, and 20 of these species have been positive for *B. ceti* or *B. pinnip-*

pedialis by culture or PCR assays (Hernández-Mora et al. 2013, Foster et al. 2015). The high seroprevalence seen here in all populations across Scotland suggests that wildlife professionals working with live seals could be exposed to the bacterium, and care should be taken when handling the animals and working with samples. To date, there have been 4 documented cases of humans infected with *B. ceti* (Brew et al. 1999, Sohn et al. 2003, McDonald et al. 2006), demonstrating the zoonotic potential of that species, but human infections with *B. pinnipedialis* have not been documented.

In conclusion, over one-quarter of Scottish harbour seals have detectable levels of antibodies to *Brucella* which may indicate endemicity in these populations, possibly to a strain of the pathogen that has little effect on the health of individuals. These prevalence rates do not appear to explain the declines in Orkney and along the East coast as the prevalence in these areas is the same as in populations along the West coast that remain stable. The causes of the decline are likely to vary between regions and are probably due to a combination of factors, but *Brucella* infection does not appear to be one of them, based on our findings and comparison with seroprevalence rates for Scottish harbour seals before 2002 (Foster et al. 2002). Despite the routine use of the serological tests used here in many assessments of exposure to *Brucella* species, further validation of the tests for marine mammals is needed, and the discrepancies between the 2 test types here highlight the need for careful interpretation of the results.

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