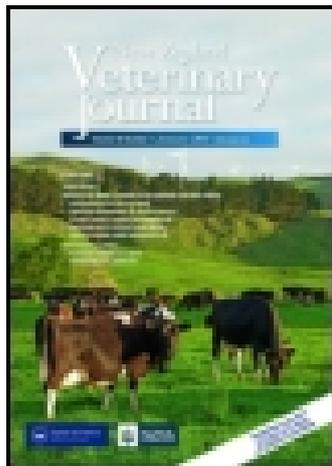


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Review Article

The epidemiology of *Mycobacterium bovis* in wild deer and feral pigs and their roles in the establishment and spread of bovine tuberculosis in New Zealand wildlife

G Nugent^{*§}, C Gortazar[†] and G Knowles[‡]

Abstract

In New Zealand, wild deer and feral pigs are assumed to be spillover hosts for *Mycobacterium bovis*, and so are not targeted in efforts aimed at locally eradicating bovine tuberculosis (TB) from possums (*Trichosurus vulpecula*), the main wildlife host. Here we review the epidemiology of TB in deer and pigs, and assess whether New Zealand's TB management programme could be undermined if these species sometimes achieve maintenance host status.

In New Zealand, TB prevalences of up to 47% have been recorded in wild deer sympatric with tuberculous possums. Patterns of lesion distribution, age-specific prevalences and behavioural observations suggest that deer become infected mainly through exposure to dead or moribund possums. TB can progress rapidly in some deer (<10%), but generalised disease is uncommon in wild deer; conversely some infected animals can survive for many years. Deer-to-deer transmission of *M. bovis* is rare, but transmission from tuberculous deer carcasses to scavengers, including possums, is likely. That creates a small spillback risk that could persist for a decade after transmission of new infection to wild deer has been halted.

Tuberculosis prevalence in New Zealand feral pigs can reach 100%. Infections in lymph nodes of the head and alimentary tract predominate, indicating that TB is mostly acquired through scavenging tuberculous carrion, particularly possums. Infection is usually well contained, and transmission between pigs is rare.

Large reductions in local possum density result in gradual declines (over 10 years) in TB prevalence among sympatric wild deer, and faster declines in feral pigs. Elimination of TB from possums (and livestock) therefore results in eventual

disappearance of TB from feral pigs and wild deer. However, the risk of spillback infection from deer to possums substantially extends the time needed to locally eradicate TB from all wildlife (compared to that which would be required to eradicate disease from possums alone), while dispersal or translocation of pigs (e.g. by hunters) creates a risk of long-distance spread of disease. The high rate at which pigs acquire *M. bovis* infection from dead possums makes them useful as sentinels for detecting TB in wildlife.

It is unlikely that wild deer and feral pigs act as maintenance hosts anywhere in New Zealand, because unrestricted year-round hunting keeps densities low, with far less aggregation than on New Zealand farms. We conclude that active management of wild deer or feral pigs is not required for local TB eradication in New Zealand.

KEY WORDS: *Tuberculosis, Mycobacterium bovis, New Zealand, wild deer, Cervus elaphus, feral pig, Sus scrofa, host status, spillover host, maintenance host, disease control*

Introduction

Eradication of tuberculosis (TB) from a multi-species wildlife host complex requires that the disease be managed in all maintenance host species. By definition, maintenance hosts are, if unmanaged, able to sustain the disease independently of other species, whereas, in the long term, spillover hosts do not. Host status is often ascribed to species as if it was a fixed characteristic, but it is not. Rather, host status is a local or regional characteristic that varies with host density and/or with variation in transmission mechanisms and other factors that affect the basic reproductive rate of the disease (Nugent 2011).

In New Zealand, introduced brushtail possums (*Trichosurus vulpecula*) are the most important wild animal maintenance host for TB (Morris and Pfeiffer 1995; Coleman and Cooke 2001; Nugent *et al.* 2015). In contrast, wild deer (predominantly red deer, *Cervus elaphus*) and feral pigs (*Sus scrofa*) are considered to be spillover hosts. Therefore, these species have not been targeted for TB management or control despite the sometimes high prevalences of TB among feral pigs, in particular, often being far higher

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LN Lymph node
TB Tuberculosis
VRA Vector Risk Area

than the prevalence among sympatric possums (Nugent 2011). However, there is strong evidence that wild boar (the conspecific progenitor of feral pigs) are maintenance hosts for TB in parts of Spain (Naranjo *et al.* 2008; Boadella *et al.* 2012), as are white-tailed deer (*Odocoileus virginianus*) in Michigan, United States of America (USA; O'Brien *et al.* 2011), red deer in parts of Europe (Gortazar *et al.* 2012), and fallow deer (*Dama dama*) on deer farms in New Zealand and elsewhere (Robinson *et al.* 1989; Griffin *et al.* 2004). Given these differences in host status, if the spillover assumption is not universally valid, the decision not to include active control of TB in wild deer and feral pigs could, potentially, undermine New Zealand's overall TB control and eradication programme. The primary aim of this review is to assess that risk and evaluate its potential importance, based on a summary of the ecology and management of these species in New Zealand and the pathology and epidemiology of TB in them.

Ecology and epidemiology of TB in wild deer and feral pigs

Status

Neither wild deer nor feral pigs are native to New Zealand; both were introduced by early European settlers in the eighteenth or nineteenth centuries and are now widespread. As in America and Europe, these animals are valued as a hunting resource (Figgins and Holland 2012) and, since the 1970s, deer have also been domesticated. There were ~1 million farmed deer New Zealand in 2012 (Anonymous 2012a). In June 2012, just five deer farms were confirmed as infected with *Mycobacterium bovis* (Anonymous 2012b), down from over 361 farms in 1987. As introduced species, wild deer and pigs are viewed by conservationists as pests due to their unwanted impacts on indigenous biodiversity (Anonymous 2001). Deliberate or incidental killing of deer and pigs as part of TB control and surveillance programmes is not, therefore, as controversial in New Zealand as it is in some other countries.

Wild deer as hosts of TB

Demography and ecology

Seven taxa of wild deer occur in New Zealand (Nugent *et al.* 2001), with a total population of ~250,000 (Nugent and Fraser 1993). Red deer are predominant (~80% of the 1988 harvest), followed by fallow deer and sika deer (*C. nippon*) (Nugent 1992a). These are the only three deer species known to have been infected with *M. bovis* in New Zealand. All three are temperate species with similar biology – an autumn mating season leading to an early summer birthing season when a single fawn is born. Females usually reproduce annually from 2 years of age, but males do not reach full maturity until 6–8 years. When not limited by food or resources, deer populations can increase by about 35% annually or higher with immigration (Forsyth *et al.* 2010). Deer can live for more than two decades, but most New Zealand populations are heavily and continuously hunted with no restrictions on age or sex of the animals harvested. As a result, the mean age of deer shot by New Zealand hunters is usually <3 years, with few males surviving to full maturity (and very few beyond 10 years) and few females beyond 15 years (Nugent and Fraser 2005).

Wild deer occupy >120,000 km² of New Zealand, which corresponds to 44% of the country (Fraser *et al.* 2000). Deer are

present in most of the 10.5 million hectares of land where TB occurs in wildlife, which are designated as Vector Risk Areas (VRA). However, the deer are mostly confined by hunting to the more heavily forested parts of the VRA, where carrying capacities are probably 15–30 deer/km² (Nugent and Fraser 2005). However, helicopter-based commercial harvesting, introduced in the 1960s, has reduced deer densities in most places, and to near-zero densities in most open areas lacking substantial cover (Challies 1991). Therefore, the overall average deer density (all species) in forests is probably <5 deer/km², with densities in the 2–5 deer/km² range in the South Island and 5–15 deer/km² in the North Island (Nugent 1992b). Densities of sika and fallow deer tend to be higher than these all-species averages, with extreme local densities of up to 40 deer/km² recorded occasionally (Nugent and Yockney 2004).

Large-scale population control of deer is feasible with the techniques available, but this is currently not part of the national TB management plan. In addition to lethal control measures, vaccination could be used to manage TB in wild deer. Inoculation with live BCG (*Bacillus Calmette-Guérin*, a non-virulent form of *M. bovis*) vaccine is effective in protecting red deer and white-tailed deer, and the availability of an effective oral formulation potentially makes it feasible to deliver the vaccine to free-ranging wild deer (Griffin *et al.* 2001; Nol *et al.* 2008). However, the lack of major impediments to lethal control makes it unlikely that vaccination of wild deer would be considered as a viable alternative in New Zealand.

Overall, the densities of wild deer in New Zealand are much lower than in places where spread of TB between deer has been confirmed such as on deer farms and hunting estates in New Zealand and overseas, where densities are typically some hundreds of deer/km² and the incidence of TB can be very high (Robinson *et al.* 1989; Griffin *et al.* 2004); or in Michigan, USA, where the disease emerged in white-tailed deer during the 1990s after deer densities increased to ~20 wild deer/km² as a result of supplemental feeding (O'Brien *et al.* 2002); or in Spain where densities of red deer averaged 19 deer/km² (range <1–66 deer/km²) across a number of areas, mostly hunting estates (Acevedo *et al.* 2008).

Year-round hunting in New Zealand has resulted in low-density deer populations that are widely dispersed with deer roaming as individuals or in small family groups. In contrast, hunting seasons overseas are typically short in duration, and outside the hunting seasons deer are frequently able to aggregate into larger herds. This natural aggregation can be intensified by supplemental feeding and at water supplies (both natural and artificial), resulting in increased interactions between deer and an increased likelihood of TB transmission (Vicente *et al.* 2007a). In Michigan, supplemental feeding has been considered the crucial factor for TB emergence in wild white-tailed deer there, but not elsewhere in the USA (Miller *et al.* 2003).

Home range size and dispersal distances help determine the scale over which deer could spread TB. In New Zealand, farm escapes or illegal liberations (Fraser *et al.* 2000) are a potential mechanism for long-distance spread of TB. Overseas and New Zealand data indicate that, where cover and food are adequate, female red deer occupy ranges of a few km², while the ranges of males tend to be substantially larger (Nugent *et al.* 2003a). In mixed forest in the West coast region of the South Island the home range sizes of six young female red deer varied between 1.7 km² and 3.8 km², but were much larger for two males at 21 and 26

km² (Nugent *et al.* 2003a). In unforested high country in New Zealand, adult male ranges can be up to 190 km² (Knowles 1997; Yockney *et al.* 2013). Fallow deer ranges are roughly half as large as that of red deer, and sika deer are intermediate (Nugent *et al.* 2001). Few female red deer disperse more than 4 km, and although most males do disperse, only 5% move more than 10 km (Nugent *et al.* 2003a).

TB origins and source

Tuberculosis was first recorded in wild deer in New Zealand in 1954 on the West Coast (Livingstone *et al.* forthcoming) and was widespread by the 1980s in both wild and farmed deer (de Lisle and Havill 1985; Lugton *et al.* 1998), with infections reported from red, sika and fallow deer (Cooke *et al.* 1999). The susceptibility of deer to TB varies widely among individuals and is influenced by climate, nutrition, age and genetic factors (Griffin and Buchan 1994; Griffin and Mackintosh 2000). Unlike cattle, farmed deer in New Zealand rarely exhibit clinical signs of the disease before the last 1–2 weeks of life (Beatson 1985; Griffin and Buchan 1994).

Before the 1960s, TB was not widespread in New Zealand wildlife, but deer probably became infected through occasional direct contact with *M. bovis*-infected cattle, or indirectly via shared feeding areas. For example, there are reports from the mid-1900s that New Zealand farmers would sometimes capture wild fawns, raise them on milk from *M. bovis*-infected dairy herds, then release them back into the wild, potentially (but unknowingly) creating a reservoir of wildlife infection (P. Livingstone¹ pers. comm.). In Michigan, USA, occasional outbreaks in cattle herds continue to occur as a consequence of TB transmission from wild white-tailed deer populations (O'Brien *et al.* 2011), and indirect deer-to-cattle transmission via shared feedstuffs has been demonstrated (Palmer *et al.* 2000). Wild deer originally infected by farmed cattle may have been the indirect conduit by which TB established in New Zealand's possum population in the 1960s (Morris and Pfeiffer 1995). After the 1960s, however, commercial aerial hunting reduced deer densities to much lower levels (Nugent *et al.* 2001), especially in open grassland areas; therefore, the habitat overlap between cattle and wild deer is now minimal and the likelihood of TB transmission between these species is low.

TB Prevalence

Before the 1990s, overseas reports of TB in wild deer populations indicated prevalences of below 6% (Clifton-Hadley and Wilesmith 1991). Since then, much higher prevalences have been reported among red deer in Spain with an average of 14%, range 0–66%, reported from 21 hunting estates (Vicente *et al.* 2006). High local prevalences have also been reported in France (up to 24% among red deer in the 2005–2006 hunting season; Zanella *et al.* 2008), and among fallow deer in England (18.5% prevalence among 65 deer examined between 1971 and 1996; Delahay *et al.* 2002). Observations during the 1980s of high prevalences of TB in farmed and wild deer in New Zealand led to speculation that wild deer could be maintenance hosts of TB (Morris and Pfeiffer 1995). However, there were early indications that TB transmission between wild deer was rare, specifically that wild fawns were rarely infected even when a high proportion of their mothers were (Nugent and Lugton 1995; Lugton *et al.* 1998). The absence of infection in fawns, however, does not

reflect resistance to infection, which was demonstrated when Palmer *et al.* (2002) induced severe infection in young white-tailed deer fawns by feeding them with TB-contaminated milk. Furthermore, in a large New Zealand outbreak in farmed red deer, over 95% of 5-month-old fawns were reported to be infected (Griffin *et al.* 2004).

There have been no longitudinal studies of *M. bovis* infection in New Zealand deer, but between 1993 and 2003 periodic cross-sectional surveys of TB prevalence were conducted in the Hauhungaroa and western Kaimanawa Ranges in the central North Island, and in the Hochstetter and Omoto Ranges in the West Coast region of the South Island (Nugent 2005). In those studies, the overall TB prevalence was 8–37% in areas where deer were sympatric with uncontrolled tuberculous possum populations; prevalence increased steadily with age for about 3 years after independence, and the increase was faster in males. This suggests that either the incidence of new infection is more or less constant with little loss of infection (through TB-induced mortality or resolution of infection) or, less likely, that the risk and incidence somehow increases with age. After about 4 years of age however, age-specific prevalence stabilised and then appeared to decline in old age, with the decline thought to reflect a combination of TB-induced mortality and some loss of detectable infection through the resolution of overt disease (Nugent 2005); consistent with that, a lower proportion of infected adult deer had typical TB lesions compared with sub-adult deer. An initial increase in the prevalence of TB-like lesions with age has also been reported among red deer in Spain (Vicente *et al.* 2006), and in an outbreak among red deer in a small (91 km²) isolated forest area in France (Zanella *et al.* 2008). In the French study, the TB prevalence in juveniles remained constant (11% and 14%) between 2002 and 2005, while that in adults increased from 13% to 32%.

In New Zealand, the prevalence of infection is similar in male and female deer, with a higher prevalence in young males (≤ 2 years of age) offset by a lower prevalence in older males (> 5 years; Nugent 2005). In contrast, in Michigan, USA, male white-tailed deer ≥ 2 years of age were more likely to be infected than females of similar age (O'Brien *et al.* 2002). Similarly, in Spain, the prevalences of infection in sub-adult and adult female red deer were found to be lower than the prevalences for males of the same age class (Vicente *et al.* 2006, 2007b).

As part of the present review, we obtained and collated unpublished historical statistics from databases held by the Ministry for Primary Industries (or their predecessors), and TBfree New Zealand, on the numbers of commercially harvested wild deer and feral pigs diagnosed as tuberculous during post-mortem meat inspection, or from formal surveys of local TB prevalence in wildlife. The deer records are presented here and the pig data in the pig section below. These records show that from mid 1994 to mid 1997, 209 tuberculous wild deer were identified among a total of 80,457 inspected (0.26% prevalence overall). In contrast, for the period mid-2008 to mid-2012, there were just 26 tuberculous wild deer detected among 74,701 inspected (0.03% prevalence), although this rate still exceeds that in livestock (which was 0.004% of 4.6 m farmed cattle and deer tested between July 2011 and June 2012; TBfree New Zealand Ltd, unpublished data). All 26 TB-positive wild deer identified between 2008 and 2012 were from north Canterbury or the West Coast regions in the South Island, where there are substantial areas in which possum control has not yet been imposed. By

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contrast, in areas subject to intensive possum control, none of 872 wild deer from 10 formal wildlife TB surveys conducted between mid 2008 and mid 2012 (mostly in the southern and central North Island) were confirmed infected, although eight animals had gross but culture-negative lesions suggestive of TB (Anonymous 2012b).

The impact of possum control on TB prevalence in wild deer is most evident in a western central part of Hauhungaroa Range in the North Island. It was here that the highest prevalence ever recorded in wild New Zealand red deer (20/44, 45%) was measured in 1999 (Nugent 2005), just before possum control was imposed for the first time. By 2011, after a decade of possum control, prevalence had declined to 0% (0/37) (Nugent et al. 2012).

Lethality

Occasional observations by researchers and New Zealand hunters of young wild deer with that are too weak to flee and have been assumed to be tuberculous, suggest that TB progresses rapidly to death in some wild deer. Infected fawns on deer farms can also die before 12 months of age (Beatson et al. 1984). However, there is wide genetic variation in susceptibility to disease in farmed red deer (Mackintosh et al. 2000), with only a few (~5–10%) appearing to be highly susceptible (Griffin and Mackintosh 2000). The same appears to be true of wild deer, with indications that some deer can survive for many years with *M. bovis* infection. This was inferred from areas where possum control had reduced the incidence of new infections in young deer to low levels (and eventually to zero), but the prevalence of infection in adult females remained unchanged, suggesting there is little loss of infection from either additional TB-induced mortality or from resolution (Figure 1). As noted by Buchan and Griffin (1990), this parallels the long latent phase post-infection followed by potential reactivation late in life that is typical of *M. bovis* infection in humans (Barry et al. 2009). The crucial consequence of deer carrying a latent *M. bovis* infection is the risk that they can act as a temporal vector, potentially carrying disease through time for a decade or more.

Pathology and lesion distribution

Although gross tuberculous lesions in deer vary in shape and form, the most frequent presentation is well-encapsulated abscesses that

are detected by lymph node (LN) enlargement; the LN often contain several granulomas. The material within lesions can be liquefactive, caseous or, particularly in older deer, highly calcified (to the point of being bone-like).

In contrast to a study in Spain, in which 57% of red deer with TB lesions presented with generalised disease and the percentage of generalised cases increased with age (Vicente et al. 2007b), generalised TB is comparatively uncommon in wild deer in New Zealand. In the central North Island, about a quarter of *M. bovis*-infected deer had no visible lesions, compared to two thirds of deer from two areas of the West Coast region in South Island (Nugent 2005). In those surveys, the deer with one or more visible lesions usually presented with a single lesion <1 cm in diameter, and many of these lesions were in the head of the animals, in either the tonsils or the retropharyngeal, parotid or submandibular LN. Of 42 deer with only a single gross lesion, Nugent (2005) reported that 42% had infection in the retropharyngeal LN of the head, compared to 15% in the mesenteric LN of the alimentary tract, 10 and 6% in the lung-associated mediastinal and bronchial LN respectively, and 6% in the popliteal LN of the hind leg. If, as is often assumed, the presence of a single lesion usually reflects a relatively early stage of infection, this lesion distribution suggests wild New Zealand deer acquire TB by multiple routes (ingestion, inhalation, and perhaps also occasionally via skin breakage). Lesion distribution resulting from intra-specific transmission among New Zealand farmed deer and in wild white-tailed deer in Michigan, USA (Hathaway et al. 1994; O'Brien et al. 2001) is broadly similar to that recorded in wild deer in New Zealand, but with lesser involvement of the mesenteric LN. In contrast, in a recent TB outbreak in red deer in France (Zanella et al. 2008), there was an unusual predominance of mesenteric lesions thought to reflect ingestion of *M. bovis* during investigation by deer of offal from hunter-killed tuberculous deer and/or wild boar. However, in Spain a high prevalence of abdominal TB lesions has also been reported among wild red deer, e.g. 54% and 41% of *M. bovis*-infected red and fallow deer, respectively (Martín-Hernando et al. 2010), and in such cases alimentary tract infection is thought to more likely reflect transmission by food and/or water.

Transmission to deer

In New Zealand, the absence of *M. bovis* infection among wild red deer fawns suggests that deer-to-deer transmission is rare. Furthermore, as fawns share the same foods as their mothers from soon after birth, indirect transmission from other host species via feed-stuffs, water, or other environmental contamination must also occur rarely. Deer aside, the other common potential sources of infection for wild deer in New Zealand are from cattle, ferrets (*Mustela furo*), pigs and possums. However, it is known that wild deer become infected in “deep” forest where they have little opportunity to interact with either ferrets or livestock (Nugent 2005), which makes it unlikely that those species are major sources of infection in all cases; and, as explained in the next section of this review, pigs are regarded as end-hosts of TB in New Zealand unless their carcasses are scavenged, reducing their likelihood of transmission of infection to wild deer. Therefore, the inference is that most of the TB observed in wild deer in New Zealand is acquired from possums. This is considered to result from deer aggressively investigating terminally ill possums (Sauter and Morris 1995).

Infection via the palatine tonsils in the buccal cavity, which drain into the retropharyngeal LN, is considered the primary route of

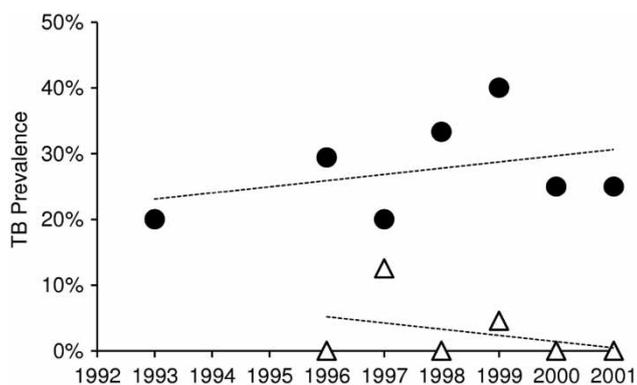


Figure 1. Trends in the prevalence of *Mycobacterium bovis* infection in female deer between 1993 and 2001 in central North Island areas of New Zealand. The upper series (circles) comprises all annual cohorts of adult and sub-adult females that were born before 1993 and exposed to uncontrolled *M. bovis*-infected possum populations. The lower series (triangles) comprises adult and sub-adult females born after the implementation of possum control measures in 1993. Adapted from Nugent (2005)

transmission to both wild and farmed deer in New Zealand (Lugton *et al.* 1998). Supporting this, experimental inoculation of the tonsils of captive red deer with virulent *M. bovis* produces a pattern of TB that is clinically similar to that seen in naturally infected farmed deer (Mackintosh and Griffin 1994). Infection through inhalation of infectious aerosols appears to be of minor importance, because lung lesions are uncommon in wild deer in New Zealand. Furthermore, historically, coughing was rarely seen in infected farmed deer, whereas in cattle, where thoracic lesions are more common, it is a common feature of the disease (Beatson 1985).

Tonsillar infection may occur when the animal is eating, drinking, licking, muzzling or grooming (Mackintosh and Griffin 1994). We consider that the predominance of infection in the LN of the head, but lesser prevalences in the LN of the thorax and of the alimentary tract, implicates licking and muzzling of some source of infection, such as dead or dying tuberculous possums, as the predominant mechanism of transmission of TB to wild deer in New Zealand, rather than ingestion with food or inhalation of airborne aerosols. Similar patterns of lesion distribution are observed among deer from countries without possums, which suggests that some form of infection involving the oral cavity can occur intra-specifically, either by licking of shared food supplies (Palmer *et al.* 2004) or by visiting communal water sources (Vicente *et al.* 2007b), or perhaps during interactions such as nose-to-nose muzzling that occur frequently at such shared sites.

Transmission from deer

Deer can excrete *M. bovis*, but in 58 naturally infected wild deer in New Zealand, excretion was minimal with viable *M. bovis* bacilli isolated from only 8% of oropharyngeal swabs, 2% of tracheal and nasal swabs, 2% of faecal samples and 0% of urine samples (Lugton 1997). Most of the positive samples in the study came from four deer that had generalised disease. The higher percentage of positive oropharyngeal swabs was interpreted as evidence that excretion via the tonsils, and therefore in saliva, may be the most common route of excretion (Lugton 1997).

On farms, a few infected deer appear to become highly infectious to other deer through development of severe disease, often involving open sinuses from peripheral lymph nodes that drain bacilli directly onto the skin (Lugton 1997; Griffin and Mackintosh 2000). Rapid spread of TB among farmed deer appears to always involve such severely infected individuals and probably occurs through direct contact with bacilli-laden discharges (Lugton *et al.* 1998). Pseudo-vertical transmission is another potential pathway for intraspecific transmission, as fawns have been experimentally infected by feeding them with contaminated milk (Palmer *et al.* 2002). However, no cases of gross lesions in mammary tissue have been reported from wild deer in New Zealand.

As there is little evidence of either direct or indirect transmission between wild New Zealand deer, indirect transmission from wild deer to farmed deer, cattle or possums is presumably also uncommon. However, cattle have occasionally acquired infection from farmed deer (Hennessey 1986). Transmission apparently occurred via pasture contamination in three cases and across a fence in another case. In all four cases, there was a high prevalence of infection in the sympatric or adjacent deer herds.

The main route of interspecific transmission from infected wild deer is likely to be other animals scavenging their carcasses;

predominantly pigs, ferrets, and possums. Pigs have been filmed feeding on deer carcasses (Nugent 2005) and ferrets on pig carcasses (Yockney *et al.* 2008), so the latter would presumably also scavenge deer carcasses. Importantly, possums occasionally scavenge meat and carcasses (Ragg *et al.* 2000) and have been filmed feeding on pig and deer remains on several occasions, albeit usually only briefly (Nugent 2005; Nugent *et al.* 2006). Such feeding has been suggested as the route by which possums first became infected in the 1960s (Nugent 2011). The presence of a single large mesenteric lesion in a possum trapped at a site where tuberculous deer carcasses had been left several months previously (Nugent *et al.* 2003a) provides some circumstantial support for this possibility. Furthermore, the detection of new *M. bovis* infection in possums within 2 years of *M. bovis*-infected farmed deer of South Island origin being moved to a previously TB-free North Island area (Mackereth 1993) provides compelling evidence of TB transmission from deer to possums.

Overall, wild New Zealand deer are most likely to transmit *M. bovis* to scavenging wildlife, and only rarely to other herbivores (including livestock). Of the potential scavengers, Nugent (2005) has suggested that individual pigs have the greatest likelihood of becoming infected from feeding on tuberculous deer carcasses simply because they consume much more of the carcass per encounter than small scavengers such as ferrets, although the latter in turn consume far more than occasional scavengers such as possums (Nugent 2005). However, the rate of *M. bovis* transmission from deer to scavengers will depend on the relative density and susceptibility to infection of the particular scavengers. Hence, although the likelihood of an individual possum actively scavenging from a tuberculous deer carcass is probably lower than the commensurate risk of pigs or ferrets doing so, the much higher densities of possums (than pigs or ferrets) in typical forested deer habitat, and their seemingly high susceptibility to low doses of *M. bovis* (Nugent *et al.*, 2013), greatly offset this lower risk.

Host status

Intraspecific transmission of *M. bovis* has been demonstrated between experimentally infected and uninfected, captive white-tailed deer, both when they were in direct contact, and when they were not in direct contact but were sharing the same housing facilities (Palmer *et al.* 2001). Transmission has also been recorded between experimentally infected red deer and uninfected red deer in a paddock (Mackintosh and Griffin 1994). As already noted, however, there is circumstantial evidence from fawns that deer-to-deer transmission is rare in the wild in New Zealand. In an experiment aimed at determining the host status of wild deer in New Zealand, trends in the prevalence of TB in deer were compared between two areas in which possums (but not deer or pigs) were controlled (Nugent 2005). TB prevalence did not decline significantly in areas where possums were not controlled, but declined to near zero levels, over 8 years, in areas where possums were controlled (Figure 2). This outcome strongly suggests that deer are spillover rather than maintenance hosts in New Zealand, and that most, if not all, infection observed in deer was being acquired from possums in the Nugent (2005) study. The same study also showed that, at a local scale (200–300 ha), the prevalence of TB in deer was strongly and positively related to the local presence of TB in possums, with the simple presence or absence of infection in sympatric possums being a stronger predictor than actual prevalence.

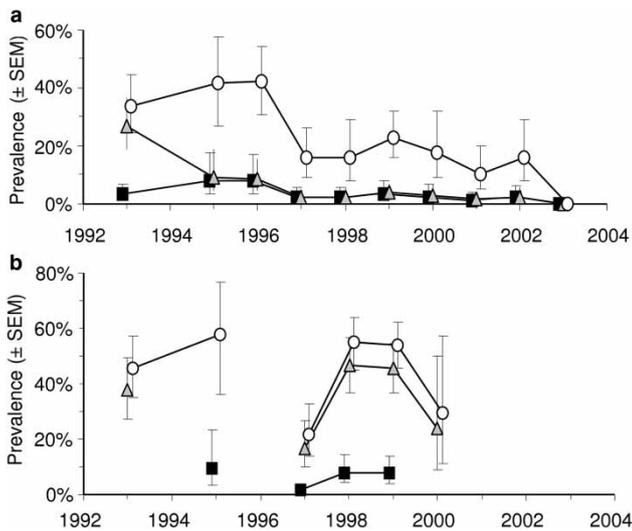


Figure 2. Mean prevalence (\pm SEM) of *Mycobacterium bovis* infection in deer killed and necropsied in the Hauhungaroa Range of the central North island of New Zealand from 1993–2003. (a) Data from the eastern side of the range where intensive possum control was applied for the first time in 1994 and (b) Data from the contiguous western side of the range where possums were not controlled during this time. Squares=0–1 year-olds, triangles=1–2 year-olds, circles=>2 year olds. Adapted from Nugent (2005)

In contrast to the New Zealand situation, the persistence of TB in white-tailed deer in Michigan, USA, for 20 years leaves no doubt that deer there are true maintenance hosts. That appears to reflect a combination of generally higher deer densities than in New Zealand, particularly in the most heavily affected areas, and far greater aggregation of deer resulting from the presence of artificial feeding sites. The threshold density for TB persistence there (in conjunction with reduced but continued supplemental feeding) appears to be ~ 12 deer/km² (O'Brien *et al.* 2011). In Spain and France, the situation is complicated by the frequent co-occurrence of red deer and wild boar, especially on hunting estates, where the TB prevalences in both species are closely correlated (Vicente *et al.* 2006). A major reduction in deer density at Brotonne, France, resulted in an eventual reduction in TB prevalence in wild red deer (Hars *et al.* 2010), implying maintenance host status in that case, but wild boar densities were also reduced so the cause and the effect in this situation are not entirely clear. In Spain, reduction in wild boar numbers in a 30 km² hunting estate resulted in modest decreases in TB prevalence among a high density (30–80 deer/km²) sympatric fallow deer herd (García-Jiménez *et al.* 2013), suggesting that the deer were acting primarily as spillover hosts, even at those high densities. Similarly, in another Spanish study, reductions in wild boar density were inferred to have resulted in reduced infection in sympatric red deer (Boadella *et al.* 2012). As transmission from infected deer to feral pigs/wild boar through consumption of carcasses (or the carcass remnants of hunter-killed deer) seems inevitable, and these two Spanish studies indicate some form of reverse transmission from wild boar to deer is possible, it is hypothesised that the two species may co-amplify the infection in each other.

Feral pigs as hosts of TB

Demography and ecology

Feral pigs are widely, but patchily, distributed in New Zealand. They occupied $\sim 34\%$ (93,000 km²) of New Zealand in 1996 after having substantially expanded their distributions in Otago,

Southland and on the West Coast of the South Island over the preceding decades (Fraser *et al.* 2000). These expansions stemmed partly from natural spread, but mostly from illegal liberations to facilitate recreational hunting.

Pigs occupy a diverse range of habitats, including native forest, plantations, rough farmland, and tussock and shrub land. They require protein-rich foods for successful reproduction and are omnivorous, eating fruits, roots, invertebrates, small vertebrates, and carrion. Possum carrion comprised 15% of pig diets in native forest in the central North Island (Thomson and Challies 1988). Pigs often feed in family groups; Coleman *et al.* (2005) filmed an adult sow eating virtually all of a possum carcass, and later returning with a litter of five 1–2-month-old piglets that all investigated the litter for remains. As *M. bovis*-infected possum carcasses carry a large burden of infective material, a single-infected possum carcass clearly has the potential to infect whole groups of scavenging pigs, which would amplify the numbers of infected animals in an area.

Extremely high pig densities were recorded in the mid 1900s in New Zealand with 123 pigs/km² (Wodzicki 1950), but such high densities have not been reported since (McIlroy 2005). Densities of 3–8 pigs/km² and 12–43 pigs/km² were recorded in heavily hunted and un-hunted areas, respectively, near Murchison, South Island in the 1980s (McIlroy 1989). Densities are maintained at these reduced levels by year-round, ground-based, recreational shooting while hunting other big game or, more commonly, by using dog teams trained to trail and corner pigs. In open areas, helicopter-based shooting was historically used to harvest pigs commercially, to control numbers, and/or collect samples for TB surveillance. Because feral pigs are gregarious, deliberately released “Judas” pigs have also been used to help locate resident pigs for TB surveillance in management or research studies (Knowles 1994; Yockney and Nugent 2006). This involves capturing or rearing feral pigs, fitting them with a radio-transmitting collars and releasing them into the wild. Radio telemetry is then used to relocate them on a number of subsequent occasions, and any resident uncollared pigs that they have become associated with are shot. This Judas method facilitates location and shooting of feral pigs at approximately twice the rate of untargeted hunting (Nugent *et al.*, 2014). With the available range of hunting techniques, it is feasible to maintain feral pig populations at very low densities in most areas, but deliberate control of feral pig populations has not been applied for TB management purposes in New Zealand. Furthermore, preventive vaccination of feral pigs has not been tested as a control measure in New Zealand, although this is being investigated overseas using either BCG or preparations of killed *M. bovis* bacilli as potential oral-delivery vaccines (Garrido *et al.* 2011; Beltrán-Beck *et al.* 2012).

There are usually no restrictions on the numbers, age or sex of pigs harvested, at least on public conservation land, and the continual hunting pressure generally prevents pigs aggregating into large groups. On some private lands and plantation forests, pigs are managed for hunting more actively though the use of closed seasons, and pig numbers are supplemented with regular releases of animals raised in captivity. The latter are usually TB-tested before release and necropsied on recovery, which helps to confirm that TB is absent from wildlife in those particular areas.

Nationwide, extrapolation from the estimated harvest of 100,000 pigs in 1988 (Nugent 1992a) using an exponential rate of increase

of 60% (see below) suggests a population size of ~110,000 at that time, which corresponds to ~1 pig/km² across the entire area occupied by feral pigs. These New Zealand densities appear to be 1–2 orders of magnitude lower than the densities of up to 90 pigs/km² recorded for *M. bovis*-infected boar populations in supplementally fed and watered hunting estates in Spain (Acevedo *et al.* 2007).

Feral pigs have a high potential reproductive rate, being able to produce up to three litters over 14–16 months, with a mean litter size of 6.2 (range 1–11, Dzieciolowski *et al.* 1992). The large litter size results in potentially larger family groups than are seen for deer. Survival of piglets is extremely variable, exceeding 90% when conditions are good, but <10% when they are poor (McIlroy 2005). Pigs have a shorter life span than wild deer, particularly in heavily hunted populations, where the mean age at harvest can be less than 1 year (Dzieciolowski and Clarke 1989). With high but extremely variable birth and death rates, the exponential rate of population increase can vary widely. For example, fluctuations between 29–118% per annum have been reported from Australia (Choquenot *et al.* 1996), which would result in booms and busts in pig numbers over time.

Adult feral pigs in New Zealand appear relatively sedentary when food and cover are adequate (McIlroy 1989), but may be more nomadic in open areas (Knowles 1994). In five New Zealand studies in largely forested areas, average home range size varied between 70 and 1170 ha (Nugent *et al.* 2003b). Knowles (1994) recorded much larger ranges (up to 23,410 ha) among 14 pigs in open mountain land.

There are few data on pig dispersal in New Zealand. In a piglet movement study at Mt White, Canterbury, no long-distance dispersal occurred, home range size ranged from 0.1–14.9 km², and the maximum distance between successive daily locations of any juvenile was 5.8 km (Barber 2004). In contrast, Knowles (1994) recorded much larger movements (up to 30 km) by individual pigs in largely unforested landscapes in central Otago. Nugent *et al.* (2002) studied 16 pigs deliberately released (as uninfected sentinels) into a forested area on the West Coast; within 6 months all were relocated and killed, by which time all but one were recovered within 13 km of their release points (the one exception was a female that had shifted 35 km). After 6 months of natural TB exposure, all 16 of these pigs had become infected with *M. bovis*, and the pattern of lesion development indicated that they had probably acquired infection within their first 2 months of release, suggesting, consequently, that the female pig that had shifted 35 km had acquired infection before she dispersed. Overall, pig home range sizes and dispersal distances are usually, but not always, larger than those of deer, particularly in habitats with limited cover.

TB origins and source

In New Zealand and globally, TB was historically common in domestic pigs, often as a result of feeding milk or offal from infected cattle (Nuttall 1986; Cousins 2001). However, TB was not identified in feral pigs in New Zealand until 1964 (Ekdahl *et al.* 1970). Since then, it has been reported in feral pigs from most areas where TB is endemic in wildlife (de Lisle 1994). Furthermore, there are no areas where the prevalence of TB is known to be high in feral pigs, but low or zero in other sympatric or nearby hosts.

Prevalence of TB in feral pigs

The first formal surveys in New Zealand quickly identified high prevalences of TB in some feral pig populations; for example, 31–32% recorded in central Otago between 1989 and 1993 (Wakelin and Churchman 1991; Knowles 1994). The infected animals were clustered in space, with up to 100% prevalence recorded within some family groups. Subsequent cross-sectional surveys of pigs in areas where tuberculous possums were present reported total-sample prevalences of 42–78%, with 100% of some age-classes infected in some areas (Nugent *et al.* 2011a).

The Ministry for Primary Industries and TBfree New Zealand databases on commercial pig and deer harvest and survey data, as referred to for deer above, indicate that, at a national scale, overall TB prevalence is low among feral pigs in New Zealand; of 12,698 feral pigs harvested commercially in the 3 years to mid 1997, only 133 (1.0%) were tuberculous. Since 2000, commercial harvesting of feral pigs has largely ceased so there are no comparable recent figures. However, in local-area surveys conducted in the 5 years to mid 2012 involving some 13,000 feral pigs, the TB prevalence was 0.2% ($n=7,245$) in areas long subject to intensive possum control, compared to 0.4% ($n=5,645$) in less intensively managed areas (TBfree New Zealand Ltd, unpublished data).

In several other countries, TB has been found in feral pigs in areas where the disease occurs in other hosts. In Australia, Corner *et al.* (1981) documented high prevalences of TB (48–85%) in feral pigs that shared habitat with tuberculous cattle (*Bos taurus*) and water buffalo (*Bubalus bubalis*). On Molokai, Hawaii, a 20% prevalence of TB was recorded in feral pigs in 1980, and TB had previously been found in both cattle and wild axis deer (*Axis axis*) (Essey *et al.* 1981). In semi-domestic, free-ranging black pigs in Sicily, *M. bovis* was isolated from 3% of pigs within an area of 1260 km², and from 27% of pigs in two infected herds of ~3000 animals each (Di Marco *et al.* 2012).

Tuberculosis has become increasingly common in wild boar throughout much of Europe, most notably in Spain, Portugal, France and Italy, but also in England and at low levels throughout much of Eastern Europe (Gortazar *et al.* 2012). In Spain, Portugal, and France, this increasing prevalence is attributed to characteristics of Mediterranean habitats, such as dry summers causing animals to aggregate at the few remaining watering sites. However, changing patterns of land use are also implicated, particularly the development of hunting estates where wild boar are maintained at high densities and managed in ways that promote aggregation at feeders and waterholes (Santos *et al.* 2012). On such estates, and a few natural areas where wild boar are protected, the prevalence of TB-like lesions is frequently higher than 50% and can reach 100% (Vicente *et al.* 2006).

In New Zealand, no *M. bovis* infection has yet been found in feral piglets that are less than 2-months of age, which suggests little or no vertical or pseudovertical transmission; after 2 months, however, TB prevalence increases with age (Figure 3). Like deer, this increase suggests an accumulation of infection over time, i.e. pigs acquire infection far more rapidly than they die of TB or resolve infection. In the northern high country of the South Island, the incidence of infection in young pigs exceeded three new cases per year of pig exposure (Nugent and Whitford 2008a). In some parts of that area, *M. bovis* prevalence approached 100% in 1–2-year-old pigs, but was lower in older

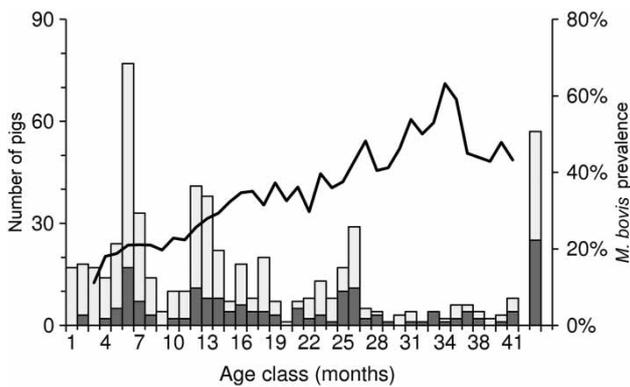


Figure 3. Five-month running average prevalence of *Mycobacterium bovis* infection (line) and sample size (columns) for 592 pigs aged 1–41 months that were necropsied between 1994 and 2008 in three regions of New Zealand. The far right bar represents all adult pigs >3.5 years old (~10% of the sample). The dark shading represents the number of tuberculosis-positive animals. Adapted from data presented in Nugent and Whitford (2008b), with permission

pigs (Nugent *et al.* 2011a). That decline cannot be attributed to better survival of uninfected sub-adult pigs as there appears to have been none; the implication is, rather, that some pigs are able to resolve detectable infection.

In studies of European wild boar, some authors reported increased prevalence rates with age (Vicente *et al.* 2006), but others reported higher prevalence rates and more generalised and severe infection in juveniles (Gortázar *et al.* 2008; Santos *et al.* 2009). The latter could reflect greater TB-induced mortality in young pigs where a combination of factors such as greater genetic susceptibility, high density and inadequate food and/or water places the population under greater stress than experienced by most feral pig populations in New Zealand. Prevalence is usually similar between the sexes, but in one Portuguese study prevalence was much higher in females (Santos *et al.* 2009).

Pathology and lesion distribution

Of all the domestic livestock species, the pig is considered to be the most susceptible to infection with *M. bovis* (Francis 1958); however, there may be some variation in susceptibility. For instance in wild boar, the progenitor of pigs, a high level of genetic heterozygosity is associated with reduced susceptibility to infection and slower progression of disease in infected individuals (Acevedo-Whitehouse *et al.* 2005). Young domestic and feral pigs rapidly develop severe lesions in the LN of the head, lungs and abdominal organs, but with time progressive fibrosis and calcification occur (Lugton 1997). Active lesions may even regress and resolve into fibrous tissue, with few or no detectable bacilli present (Ray *et al.* 1972; Bollo *et al.* 2000). In general, numbers of viable bacilli in lesions in feral pigs in New Zealand appear to be low, especially compared to those in possums and ferrets (Cooke *et al.* 1999). In a North Canterbury study, samples from 15/43 (35%) of >3-year-old feral pigs with suspect TB lesions were negative on first culture (Nugent *et al.* 2011a). The culture negative samples were typically from lesions that were either heavily calcified or were fibrotic with few or no necrotic foci. However, for a sub-sample these ($n = 10$), five proved positive after a second re-culture, indicating that low numbers of viable bacilli were present that were on the borderline of detectability using mycobacterial culture methods.

In New Zealand, almost all pigs with visible lesions have infection in the sub-maxillary LN, and those without head lesions usually have gross lesions in the alimentary tract (Wakelin and Churchman 1991; Lugton 1997; Nugent *et al.* 2011a). This distribution is likely to be a consequence of pigs ingesting tuberculous carrion. The above-mentioned studies show that generalised disease affecting many body regions is usually rare in New Zealand. By contrast, in a Spanish study of wild boar, LN in the head were the most frequently affected (107/116 animals, 92%) but, unlike New Zealand observations, 38% of boar also had TB lesions in the lungs (Martín-Hernando *et al.* 2007).

Lethality

Of the large mammalian hosts, pigs (particularly adults) appear to be the least detrimentally affected by *M. bovis* infection. Consistent with that, New Zealand pigs rarely show clinical signs of TB. In a study in Hochstetter Forest, West Coast region, South Island, all of 16 released pigs that became *M. bovis*-infected within 2 months of their first exposure to natural infection, were still in good condition when the animals were killed up to 21 months later, and none had external signs of disease (Nugent *et al.* 2002). The ability to tolerate infection, coupled with the accumulation of infection with age, implies that TB-induced mortality is low in feral New Zealand pigs, at least within the first year infection.

Transmission to pigs

The predominance of lesions in the head and mesenteric LN in feral pigs in New Zealand is likely to result from scavenging tuberculous carrion, such as possums (Nuttall 1986; de Lisle 1994; Lugton 1997). Ramsey and Cowan (2003) suggested that about 80% of *M. bovis*-infected possums die in places accessible to pigs. In one 400 ha South Island high country area containing over 20 pigs, pigs found and scavenged 32 of 40 possum carcasses that had been randomly placed in the area (Barber 2004). Based on broad best guesses as to pig and possum densities on Molesworth Station, Marlborough region, Nugent and Whitford (2008a) estimated that an individual uninfected pig would have a high (0.38–0.74) probability of finding and becoming *M. bovis*-infected if a tuberculous possum carcass was located within its home range.

Pseudo-vertical transmission (post-partum transmission between sows and their dependent offspring) has been suggested as an explanation for the clustering of infection in family groups of piglets (Knowles 1994; Lugton 1997). However, horizontal transmission can sometimes occur from an early age, as demonstrated by a litter of eight ~2-month old piglets killed in Hochstetter Forest that were all infected with the same *M. bovis* strain, while their mother was infected with a different strain (Nugent *et al.* 2003b). Pigs may also become infected via environmental contamination (Albiston *et al.* 1954), although such contamination is thought to be rare in New Zealand (Lugton 1997).

Deer carcasses are also readily scavenged by pigs (Nugent 2005). However, pigs appear less likely to cannibalise the carcasses of other pigs, or to feed on ferret carcasses. Cannibalism does occur (Nugent *et al.* 2003b), but in video-monitoring of 23 sets of pig remains (carcass, or viscera, or carcass with viscera exposed) in the South Island high country, Yockney and Nugent (2003) demonstrated that none were scavenged by other pigs. In a related trial, no ferret carcasses placed in pig habitat in either summer or winter were scavenged by pigs (Byrom 2004).

Transmission from pigs

Low numbers of viable bacilli in lesions, coupled with the relatively high level of organised and encapsulated infection seen in affected LN of many older pigs (Francis 1958), suggest excretion of *M. bovis* by pigs is uncommon. Correspondingly, Lugton (1997) found no bacilli in the urine, faeces or nasal cavities of four tuberculous feral pigs with widespread and large lesions. Nevertheless, there are occasional reports in New Zealand of grossly diseased feral pigs with draining abscesses (Lugton 1997). Therefore, environmental contamination may occasionally occur, and transmission to herbivores (cattle, deer, possums) via pasture is theoretically possible in New Zealand (Lugton 1997).

In Mediterranean Europe, TB often occurs in cattle herds in the vicinity of infected wild boar or semi-feral pig populations (Richomme *et al.* 2010; Vieira-Pinto *et al.* 2011; Di Marco *et al.* 2012; Rodríguez-Prieto *et al.* 2012) suggesting that transmission from pigs to cattle can occur. In contrast, in New Zealand, the main route of transmission from infected pigs to other hosts is more likely to be via carcasses of pigs that have died naturally or have been killed by hunters. Ferrets are likely to be the most at-risk species; in one trial, 10 of 13 sets of pig remains left in ferret-prone habitat were scavenged by ferrets feeding in family groups of up to five animals (Yockney and Nugent 2003). In that trial, a possum was also filmed feeding on mesenteric tissue from a pig, indicating a possible spillback mechanism from pigs to possums. Supporting that, a field study of three possum populations investigated scavenger contact using pig or deer meat containing dye-impregnated LN showed that 0–7% of the possums in each population had ingested the dye (Nugent and Whitford 2007).

Host status

Feral pigs have long been regarded as spillover or end hosts of TB in New Zealand, based on their ability to become infected without developing overt clinical disease, in contrast to most other wildlife in New Zealand (Coleman and Cooke, 2001). End-host status is supported by evidence from the Northern Territories of Australia, where TB prevalence in feral pigs declined following the removal of infected cattle and buffalo (true maintenance hosts) without any additional control of pigs (McInerney *et al.* 1995). TB transmission between live pigs can occur, at least in captivity (Ray *et al.* 1972), but only limited transmission between captive feral pigs has been observed in New Zealand, even when the animals were kept at densities far higher than those seen in the wild (Nugent *et al.* 2011b). In a formal test of the host status of feral pigs in New Zealand, the TB prevalence in such pigs fell substantially in three areas after intensive possum, but not pig, control was applied at geographic scales far greater than most pig home range sizes (Nugent *et al.* 2011a). In three other areas where neither possums nor pigs were controlled, there were only small or no declines in TB prevalence levels among the feral pigs. Furthermore, the declines in prevalence occurred rapidly after possum control was instigated, with near-zero prevalences recorded within 2–3 years in some areas. The rapidity of the decline in TB prevalence is surprising given the likelihood that some pigs can survive for longer than 2–3 years in an infected state. This suggests very high rates of population turnover due to continual hunting pressure, perhaps coupled with rapid resolution of detectable infection when pigs are not continually exposed to re-infection. The rapid decline in TB prevalence among pigs contrasts sharply with the much slower declines observed in wild deer after possum control

measures are instigated. This indicates that infected pigs are far less likely to act as temporal vectors of TB in New Zealand than infected wild deer.

The reverse experiment, in which the density of wild boar was reduced without reducing numbers of the other common TB hosts, has been conducted in Spain. The results of this work resulted in a decline in TB prevalence among the boar, suggesting boar are maintenance hosts for TB in that country (Boadella *et al.* 2012). Further evidence of maintenance host status in overseas studies is presented by the persistence of a high prevalence of TB in wild boar on hunting estates where they have been isolated from livestock and other wild ungulates (Naranjo *et al.* 2008; Gortazar *et al.* 2012). Furthermore, more frequent generalisation of infection to multiple anatomical regions was also recorded among the boar in these studies, including lesion development in the lungs (particularly in young animals). In Sicily, free-ranging domestic pigs are also considered to be maintenance hosts as there is evidence of strains of *M. bovis* being present in pigs that are not present in sympatric cattle (Di Marco *et al.* 2012). In contrast, in France, wild boar in the Brotonne Forest are considered to be primarily spillover hosts, although disease management measures are undertaken there to prevent boar from scavenging the offal of other pigs (Zanella *et al.* 2012).

In conclusion, the spillover host status of feral pigs in New Zealand is likely to be the result of comparatively low intra-specific contact rates, which in turn result from the unrestricted year-round New Zealand hunting system reducing average densities to low levels and keeping the pigs widely dispersed. In contrast, wild boar and feral pigs in Europe occur at much higher densities, and can be aggregated further by the use of artificial feeders or at watering points during hot dry Mediterranean summers, sufficient to facilitate intra-specific *M. bovis* transmission.

Roles of wild deer and feral pigs in the New Zealand TB control programme

Wild deer

Deer are now considered to be density-dependent maintenance hosts of TB, and the density threshold for disease persistence via deer-to-deer transmission alone is greater than the densities at which deer routinely occur in the wild in New Zealand. Therefore, direct management of the disease in deer (by density reduction or vaccination) is not seen as an essential prerequisite for local eradication of the disease from wildlife.

Nonetheless, the ability of deer to survive in an infected state for a decade or more imposes an important temporal constraint on the length of time believed to be required to locally eradicate TB from wildlife. This is because there is a presumed (but as yet not fully proven) risk of spillback infection from deer to possums. The strategic approach to local eradication of TB is predicated on reducing possum densities to very low levels, then ceasing control as soon as possible (to avoid on-going costs). This can be often achieved in as little as 5–10 years (Barron *et al.* 2013). But if possum control was stopped as soon as the population was deemed free of TB, possum densities could increase quickly to above the threshold density at which TB can persist indefinitely in that population, at which point spillback infection to possums (from a deer infected more than 10–15 years previously)

could therefore result in the *M. bovis* re-establishing in the possum population

Modelling this potential mechanism of long-duration spillback risk, Barron *et al.* (2013) predicted that the risk of *M. bovis* spillback from deer to possums could persist for up to 14 years after the initial reduction in possum population. The modelling also predicted that reducing deer densities by 80% would have only a small impact in reducing the duration of the risk of spillback from deer to possums. Therefore, the current possum control regime for the forested areas in which most *M. bovis*-infected deer typically reside involves applying control measures for at least 10 years, after which the possum population would take 5–10 years to return to pre-control levels. That combined period exceeds the predicted duration of any perceived spillback risk from deer, mitigating the threat of TB re-establishing in possums, but at the cost of extended possum control. The temporal constraint imposed by *M. bovis* infection in wild deer is that it largely precludes the strategic possibility of speeding up eradication from possums by applying more intensive possum control, at least in areas where wild deer and possums co-exist.

There is also a risk that deer might spread TB beyond the boundaries of the areas in which disease is already established in possums and, through spillback, establish a new focus of infection in uninfected possum populations. Infected deer have the potential to carry TB with them when they disperse, and young males can disperse for over 30 km (Nugent 2005). Spatial data on TB prevalence in deer in the Hauhungaroa Range in the central North Island indicate the scale of TB spread by deer (Nugent 2005). Possum control in the eastern half of the area in 1994 greatly reduced transmission from possums to deer in the eastern half, but not in the contiguous area immediately to the west where no possum control measures were taken. Over time an east-west prevalence gradient developed within the eastern half, with a high TB prevalence on its western boundary and a declining prevalence with increasing distance eastward from that boundary, i.e. with increasing distance from an uncontrolled possum population. In female deer, prevalence fell to near zero levels within 3 km eastward of the western boundary, compared to about 5 km in males.

The parts of New Zealand known or believed to contain tuberculous wildlife are formally designated as VRA. The boundaries of these are typically set at distances of 15–20 km beyond the known or suspected location of infected possums, creating possum management buffer zones aimed at preventing TB spread beyond the VRA. As the buffer zones are substantially wider than the gradient described above, and because the prevalence of TB in possums (and therefore deer) near the edges of these VRA is usually low, so the risk of spread of TB by deer across the buffer zones seems likely to be very low.

Beyond the risks posed by being infected with *M. bovis*, deer have some functional utility in the New Zealand TB management programme. As spillover hosts, they can be used as sentinels to help assess the distribution of infection in wildlife or, where efforts are being made to eradicate TB from possums, to help assess the likelihood that TB is absent from possums in that area. For the latter, a spatially explicit Bayesian framework has been developed for calculating the probability that possums in an area are free of TB (Anderson *et al.* 2013, forthcoming). This framework makes use of negative TB diagnoses not only from possums themselves, but also from ferrets, pigs,

and deer. Comparison of the densities and TB prevalence and incidence rates for possums, deer, pigs, and ferrets, indicates that deer are up to 98% less sensitive TB-sentinels than pigs (Nugent 2005; Nugent and Whitford 2008a). This is presumably because pigs are likely to actively seek out possum carcasses, and then consume all of the infectious material they contain. It is also often more expensive to obtain deer as surveillance samples, but deer may still act as useful sentinels in areas where there are few pigs or ferrets (Nugent and Whitford 2008a).

Feral pigs

This review re-affirms the long-standing belief that feral pigs in New Zealand are spillover hosts for TB. In contrast, wild boar in Mediterranean Europe are maintenance hosts, and this is likely to be a consequence of differences in climate and ecology, and wildlife management systems between the two regions resulting in differing average densities and degrees of aggregation. Therefore, pig control or vaccination is not considered to be essential for TB eradication in New Zealand. However, feral pigs are not always end-hosts, they do occasionally pass on infection under the following circumstances: (i) to other pigs (but at too low a rate to sustain on-going infection); (ii) to cattle and deer (through environmental contamination, albeit rarely); (iii) to ferrets (through scavenging, which may occur frequently where they share habitat); (iv) to possums (also through scavenging). This suite of possible transmission routes from pigs to other species is inconsequential in areas where TB is prevalent in possums, and any residual risk of *M. bovis* transmission from pigs will almost completely disappear within a few years of TB being eradicated from possums.

Nonetheless, pigs do pose some risks to the success of TB control measures. For instance, pigs may disperse TB widely, both naturally and with human assistance. The natural dispersal risks are much the same as noted above for deer, but the less stable home ranges of pigs and their greater propensity to disperse over long distances suggest that the “risk radius” for pigs is probably greater than that for deer. Some evidence for that comes from the Hauhungaroa Range study (Nugent 2005), where intensive possum control was initiated in the 7 km-wide eastern section in 1994, but not fully in the contiguous western section until 2005. Despite this control, the *M. bovis* prevalence in older pigs remained high throughout the 7 km-wide eastern side until 2005 (Nugent *et al.* 2011a) whereas in deer, by contrast, there were no new infections in the easternmost half of the east side. Together, these observations imply that a source of infection (tuberculous possums) persisted in the western section until at least 2005, and that the wider-ranging pigs had acquired infection there and had relocated back eastwards bearing disease (i.e. they had moved up to at least 7 km) whereas the deer had ranged <4 km and consequently remained disease-free.

Adding to the risk of geographic spread of TB posed by pigs naturally dispersing, both live pigs and pig carcasses are often moved long distances by hunters. Illegal releases of *M. bovis*-infected pigs in TB-free areas offer a plausible explanation for some TB outbreaks that have occurred outside known infected areas. This hunter-related dispersal risk appears to be greater for pigs than for deer, because pigs are usually transported with the head on the body. The head of the pig is discarded later, sometimes in areas that are freely accessible to possums and ferrets (Yockney *et al.* 2008).

Pigs, in conjunction with ferrets and deer, may also increase the cost of eradicating TB from wildlife by obscuring the fact that TB has been eradicated from possums in an area. One plausible scenario involves an infected deer dying a number of years after the disease has disappeared from possums and this initially causing new infection in young pigs and/or ferrets, followed by the infected pigs themselves subsequently dying and additionally passing the disease to ferrets, both of these actions potentially amplifying the spillback risk posed by deer. If possum numbers are kept low, that risk is likely to be negligible, but if infection is detected in pigs or ferrets, the precautionary principle may force managers to continue expensive possum control for longer than actually required.

Those risks and problems aside, pigs are likely to play an increasingly important role in TB management in New Zealand, as the strategic aim is currently to eradicate TB from wildlife over at least 2.5 million ha (25% of the VRA) by 2026 (Hutchings *et al.* 2011). Pigs are recognised as being the most cost-effective sentinels for helping to confirm TB freedom in a specific area, particularly in forested areas where direct surveillance of low-density possum populations is difficult and expensive (Nugent and Whitford 2008a). Their utility as sentinels reflects their large home range size, their propensity for scavenging possum carrion, their high susceptibility to acquiring infection and presenting detectable disease, and the ability to obtain samples at low cost in many areas by collecting pigs heads from recreational hunters. A sampling intensity of just one feral pig per km² (i.e. 3–4 pigs per pig home range) is sufficient to provide 95% confidence that TB is not present in possums over the same area (Nugent and Whitford 2008a). The high sensitivity of pigs as sentinels has supported the deliberate release of radio-tagged TB-free pigs for disease surveillance purposes (Nugent *et al.* 2002), and sometimes these pigs' movements have been tracked by radio-collaring them with GPS-transmitting devices to better define the area "surveyed" (Nugent *et al.* 2011c), and sometimes to help locate non-collared sympatric resident pigs for sampling (Yockney and Nugent 2006; Nugent *et al.*, 2014).

Concluding summary

Although deer can independently maintain TB when their densities are very high (as on farms), and while increased transmission can be facilitated by local management practices and conditions (as in Spain, France, and Michigan), there is no evidence that such high densities facilitating independent *M. bovis* maintenance occur in wild deer in New Zealand. The elimination of TB from possums (and livestock) is therefore expected to result in the eventual disappearance of the disease from wild deer, and additional deer-control measures will not be necessary for TB eradication.

However, infected deer can survive for more than a decade, which imposes a temporal constraint on the duration of time for which local possum control needs to be maintained. Deer can also disperse for long distances, which can result in the need for wide containment buffers around areas containing infected possums and sympatric deer. Fortunately, possum control is usually maintained for at least 10 years, so this constraint is more apparent than real. Modelling studies also suggest that further deer control measures would be unlikely to be a cost effective means of reducing the risk of spillback from deer to possums (Barron *et al.* 2013).

The same is largely true for feral pigs, except that their shorter life-span results in a shorter duration of the risk for local spillback to possums. The other main epidemiological risk connected to feral pigs lies in the potential for long-distance spread of *M. bovis* via the transportation of infected pigs or pig carcasses by hunters. This risk has diminished substantially as possum control has expanded to cover more than 80% of the *M. bovis*-infected areas of New Zealand.

The most important gaps in current knowledge involve a lack of understanding about both the duration of time for which the spillback risk persists in deer, and how frequently scavengers such as pigs and ferrets acquire infection from aged deer. That knowledge could be important for guiding decisions about when to stop possum control. It would be particularly useful in those situations where several years of surveillance in a given area provide a high level of confidence that *M. bovis* infection is absent from possums, but the disease is then detected in an aged deer or in a young pig or ferret.

We conclude that wild deer and feral pigs are important but play a secondary role in the complex epidemiology of TB in New Zealand. The subsidiary roles they play are largely dependent on their interaction with other hosts, particularly possums. The major future involvement of pigs and deer in the active management of TB in New Zealand will lie predominantly in their use as sentinels. However, occasional unexplained incidents of TB in deer, pigs or ferrets in areas close to being, or having recently been, declared TB-free could feasibly continue, either as a result of illegal translocation and release *M. bovis*-infected pigs or via long-term persistence of viable *M. bovis* in deer. In reality, such occurrences are rarely likely to be of epidemiological significance, but the uncertainty about local TB freedom that they cause is likely to be too great to be entirely ignored, resulting in substantial precautionary expenditure on TB control measures.

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