### **Rapid Communication**

# Drought, disease, defoliation and death: forest pathogens as agents of past vegetation change



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ABSTRACT: The temperate and boreal forests of Europe and North America have been subject to repeated pathogen (fungal disease and phytophagus insect) outbreaks over the last 100 years. Palaeoecology can, potentially, offer a long-term perspective on such disturbance episodes, providing information on their triggers, frequency and impact. Mid-Holocene declines in *Tsuga* and *Ulmus* pollen dominate the Quaternary literature on forest pathogens, yet the role of pathogens, and even the presence of pathogenic fungal diseases, during these events has yet to be established. Pathogen-focused research strategies, informed by the sequence of events documented in modern outbreaks, and undertaken at high temporal resolution using a multi-proxy approach, are required. It is argued that forest pathogens are likely to have been significant agents of past vegetation change, even in cases where climate change was the primary stress factor. Copyright © 2013 John Wiley & Sons, Ltd.

KEYWORDS: climate change; disturbance; forest pathogens; Holocene; palaeoecology.

#### Introduction

A recently emerged invasive fungal disease is currently threatening populations of common ash (Fraxinus excelsior L.) across temperate Europe. Ash dieback, first recorded in Poland in the early 1990s (Husson et al., 2011; Kraj et al., 2012), had by 2010 been reported from 22 countries (Timmermann et al., 2011). The causal agent appears to be a new species of ascomycete (Hymenoscyphus pseudoalbidus Roberge ex Desm.), a sexually reproducing form of the fungus Chalara fraxinea (T. Kowalski) (Kraj et al., 2012). Ascospores, released from fallen leaves and twigs and dispersed by wind (Timmermann et al., 2011), are thought to be the primary source of infection and responsible for the rapid spread of the disease, although the commercial trade in seedlings has also been implicated (Pautasso et al., 2013). Seedling death often occurs within a few years, while in mature individuals the disease becomes chronic and the weakened trees vulnerable to attack by secondary pathogens, such as Armillaria (Timmermann et al., 2011). The immediate outlook for ash, along with the biodiversity associated with this tree in Europe, is dismal. For example, it has been estimated that 60-90% of ash stands in Denmark are affected and likely to disappear (van Opstal, 2011).

The temperate and boreal forests of Europe and North America have been considerably altered by repeated outbreaks of fungal diseases over the last 100 years. Europe has seen the loss of alder and oak populations due to epidemics of various *Phytophthora* spp. (Jung *et al.*, 2000; Thomas *et al.*, 2002; Brasier *et al.*, 2004), while the forests of North America have experienced outbreaks of beech bark disease (*Neonectria coccinea* var. *faginata* Lohm.) and both continents have been affected by chestnut blight (*Cryphonectria parasitica* [Murr.] Barr) and Dutch elm disease (*Ophiostoma ulmi* [Buisman] Nannf. nd *O. novo-ulmi* Brasier) (Schlarbaum *et al.*, 1997; Sutherland *et al.*, 1997; Brasier and Buck, 2002). Wood-boring 'bark beetles' serve as vectors for beech bark disease and Dutch elm disease and insect larvae can also be

\*Correspondence to: M. Waller, as above. Email: m.waller@kingston.ac.uk important agents of high intensity and large-scale disturbance events through defoliation. In North America the forest tent caterpillar (*Malacosoma disstria* Hübner) infected millions of hectares of sugar maple (*Acer saccharum* Marsh.) between 2002 and 2007 (Wood *et al.*, 2010), while eruptions of the gypsy moth (*Lymantria dispar* L.), introduced from Europe in the 19th century, may extend over thousands of hectares in any one year (Liebhold *et al.*, 2012).

Today fungal diseases and phytophagus insects (hereafter referred to as 'pathogens') are a major source of disturbance in temperate and boreal forest ecosystems and influence species diversity and distribution, structure and ecosystem function (e.g. Castello *et al.*, 1995; Holdenrieder *et al.*, 2004). Long-term records could make a significant contribution to a better understanding of certain aspects of such episodes, notably their triggers, recurrence interval and enduring impact. This article details the sedimentary evidence relating to past and conjectured past outbreaks and examines both how palaeoecological research might more effectively be used to document such outbreaks and the role of pathogens as agents of past vegetation change. First, the nature and causes of recent pathogen episodes are briefly reviewed.

#### Pathogens in the present

In addition to large-scale events, the activities of pathogens are perpetual within woodlands and responsible for weakening, as well as killing, individual or small groups of trees. They tend to be, but are not always, species-specific. Other species are indirectly influenced by changes in competition. Pathogenic fungi infect root, stem and foliar tissue. Trees weakened by age, shade or drought are particularly liable to attack, although fungi may also be primary agents of mortality. Defoliation by insects weakens trees through the depletion of carbohydrate reserves. With short life cycles and high reproductive potential, insect populations can fluctuate substantially. Some persist at low population densities and periodically expand to epidemic proportions, resulting in the total defoliation of host trees. The causes of the latter have been subject to much debate, but are considered influenced by the populations of both generalist predators and speciesspecific pathogen/parasites of the insects (Dwyer *et al.*, 2004). Bark beetles comprise a diverse group of insects (Coleoptera, Curculionidae, Scolytidae) that feed largely on the phloem tissue of woody plants. Their attacks reduce growth and can be significant in causing mortality. Many interact with pathogenic fungi, targeting infected trees as well as acting as disease vectors.

A number of processes can cause infectious plant diseases, such as ash dieback, to emerge (Anderson et al., 2004; Desprez-Loustau et al., 2007). The geographical range of the pathogen can be extended. In these circumstances, the vulnerability of the host is likely to be high due to the absence of coevolution with the pathogen (Brasier and Buck, 2002). Pimentel et al. (2001) estimate that 65-85% of plant pathogens worldwide are exotic to the area where they are now found. Human activity, i.e. long-distance trade and the transport of specimens for planting, has been implicated in the spread of many plant pathogens. For example, chestnut blight is believed to have been introduced into the northeastern USA on Asian chestnut trees in the late 19th or early 20th century (Pridnya et al., 1996). A number of the more recent fungal outbreaks, including Dutch elm disease, evidence for which pre-dates the 20th century (Rackham, 2003), and ash dieback can be attributed to hybridization (Brasier, 2001; Husson et al., 2011) which may allow a pathogen to escape previously effective defence mechanisms. Increased hybridization may also be a consequence of human activity, with greater mobility resulting in the co-occurrence of related, but previously allopatric, species.

Changes in the host and/or in the environment are a further major trigger for emerging plant diseases. The genetic diversity of the host influences the impact of the pathogen. For example, resistance to Dutch elm disease in Britain seems to be greater in *Ulmus glabra* (Huds.), which reproduces sexually and is therefore genetically diverse, in comparison with the more vulnerable *U. procera* (Salisb.), which reproduces clonally and may be a genetically unaltered Roman import (Gil *et al.*, 2004). Environmental change can impact directly upon the health of the host (e.g. the effects of drought on tree physiology) or upon the pathogen through effects on development and survival. The ability of some foliar disease fungi to produce spores and infect hosts is, for example, related to temperature and precipitation (e.g. Gadgil, 1977; Woods *et al.*, 2005). There is also the potential for climate to influence the spread of fungi though a series of plant–pathogen interactions from extending the range of vectors, to changing tree resistance by controlling nutrient availability.

When considering the role of pathogens in forest ecosystems it is useful to recognize that disturbance events often consist of a series of interactions between abiotic and biotic factors. Primary factors (climate or a pathogen), responsible for initial stress, weaken trees which, with their resistance compromised, are then vulnerable to secondary factors (bark beetles, fungal diseases) which can continue attacking trees even if the primary factor is no longer active (Table 1). Death, accompanied by saprophytic colonizers, can occur many years later. The affected area, after a time lag of a number of years, may then be vulnerable to fire because of an increase in the amount of combustible material. Thomas et al. (2002) have detailed this sequence of events in episodes of oak decline in central Europe. Two primary factors are considered of greatest importance: insect defoliation and either winter frost or summer drought, although infection of foliage by powdery mildew (Microsphaera alphitoides Griff. et Maubl.) or the roots by Phytophthora spp. can also contribute to the initial stress. Of the secondary causes, wood borers of the genus Agrilus have the greatest impact on the health of the tree. Fungi (Armillaria spp.) occur on oak roots in the early stage of dying, but appear largely to be saprophytic rather than agents of decline. Considering the long-term impact of exotic pathogens on forest ecosystems, Rackham (2003) suggests that because of the introduction of powdery mildew into Europe in the early 20th century (Thomas et al., 2002), oaks are now less able to regenerate under shade.

#### Pathogens in the sedimentary record

A number of recent pathogen outbreaks have been examined in the sedimentary record, with such investigations seen as a means of testing the potential for reconstructing biotic disturbances over long time scales using palaeoecological techniques (e.g. Davis, 1981; Allison *et al.*, 1986). The first

 Table 1.
 Stages in forest disturbance events in which fungal diseases and/or insect infestations ('pathogens') are primary or secondary factors; the example of modern oak dieback and their recognition in the sedimentary record.

Stage	Oak dieback	Associated changes in pollen strati- graphy	Other potential sources of sedimentary evidence
Primary stress factor(s)	Summer drought Winter frost Powdery mildew <i>Phytophthora</i>	Rapid decline in host/and other vulnerable taxa	<i>Stress</i> : tree ring analysis <i>Causes</i> : palaeoclimatic data (stable isotopes and from other techniques), pathogens as below
Secondary factors (pathogens)	Bark beetles ( <i>Agrilus</i> )	Continued low values of host, disturbance indicators, recovery of non-host species	Macrofossils: bark beetles, defoliating insects, leaf damage, fungal bodies, wood remains Spores of parasitic fungi Ancient DNA
Tree death and colonization by saprophytes	Armillaria infection	As above	Spores of saprophytic fungi
Abundance of dead wood leading to more frequent fires			Macro/micro-charcoal data
Long-term change in forest composition	Failure of oak to regenerate under shade	Lower long-term abundance of host	
Genetic change in pathogen or host		Recovery in the abundance of host	

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such studies were undertaken from the area affected by chestnut blight, which between 1904 and 1950 spread over eastern North America killing 85% of mature Castanea dentata ([Marsh] Borkh.) trees (Anderson, 1974). The declines in Castanea pollen in lake sediments are sudden, sustained and roughly synchronous over large areas (Anderson, 1974; Davis, 1981). At some sites, increases in tree species associated with chestnut are recorded after the decline, notably Betula and Quercus (Brugam, 1978). However, attempts to examine the impact of the outbreak on forest composition more widely are hampered by other unrelated land-use changes (Patterson and Backman, 1988). The late 1970s outbreak of Dutch elm disease in southern England has also been subjected to pollen-stratigraphic investigation from a wooded area (Perry and Moore, 1987). A decline in elm pollen is followed by increases in pollen types derived from outside the woodland, indicating changes in pollen dispersal following the disturbance.

More recently, Anderson et al. (2010); Morris and Brunelle (2012) and Morris et al. (2013) have produced high-resolution stratigraphic records from small sub-alpine lakes in the western USA affected by infestations of the spruce bark beetle (Dendroctonus rufipennis Kirby) during the 20th century. Fluctuations in the pollen representation of the host tree (Picea engelmannii Parry) and species not affected (notably Abies lasiocarpa [Hook.] Nutt) correspond well, in terms of both the timing and the direction of change, with historical records. Morris and Brunelle (2012) and Morris et al. (2013) indicate that ratios of host to non-host pollen are a robust method of detecting high-magnitude events. However, changes in the abundance of shrub and herb pollen are shortlived and difficult to detect and expected increases in soil erosion and alterations in nutrient cycling during outbreaks could not be identified in the sedimentary record. Nevertheless, these authors caution that additional lines of evidence (to palynology) are required if Holocene events are to be identified with any certainty.

Two mid-Holocene pollen-stratigraphic events dominate the Quaternary literature on plant pathogens: the hemlock (*Tsuga*) decline in eastern North America and the pan-European elm (*Ulmus*) decline. Both were apparently synchronous across broad geographical regions, with age of the *Tsuga* decline determined at 60 sites to be ca 5 500 cal a BP (Bennett and Fuller, 2002) and the *Ulmus* decline having a mean date of  $5 036 \pm 247^{14}$ C a BP (ca 5 800 cal a BP) in the British Isles (Parker *et al.*, 2002). Two further unusual features are associated with these events at many, but by no means all, sites. First, the declines in pollen frequencies are of a large magnitude and rapid, with data from annually laminated sediments suggesting at some sites they occurred over a period of <20 years (Allison *et al.*, 1986; Peglar, 1993). This suggests the death of populations rather than simply the absence of recruitment (Birks, 1986). Secondly, the impact on the pollen representation of *Tsuga* and *Ulmus* is long-lasting which, it has been suggested, indicates the first outbreak of a pathogen (Davis, 1981).

Attributed by Deevey (1939) to climate change, Davis (1981) compared the mid-Holocene hemlock decline with the pollen record of the 20th century Castanea decline and, based on the similarities and the features outlined above, concluded that the decline was triggered by the outbreak of a pathogen. Subsequently, finds of the remains of defoliating insects and damaged hemlock needles coincident with the decline (Anderson et al., 1986; Bhiry and Filion, 1996) appeared to corroborate the pathogen hypothesis (Table 2). However, this was not universally accepted as the primary causal process and more recent studies using further lines of evidence, including lake level data, climate modelling and stable isotopes (Hass and McAndrews, 2000; Calcote, 2003; Shuman et al., 2004; Foster et al., 2006; Zhao et al., 2010), strongly suggest the mid-Holocene hemlock decline was triggered by abrupt climate change. Most recently, Oswald and Foster (2012) indicate the event varied geographically and that in some areas a series of droughts produced a stepped decline in Tsuga pollen.

The elm decline was initially attributed to climate change, increasing continentality (e.g. Iversen, 1944), then human activity (Troels-Smith, 1960). However, over the last 30 years a broad consensus has emerged that such a widespread and complex event cannot be explained by a single process and that a pathogen is likely to have been a contributory factor (e.g. Birks, 1986; Girling, 1988; Peglar, 1993; Peglar and Birks, 1993; Parker et al., 2002; Rackham, 2003; Clark and Edwards, 2004; Lamb and Thompson, 2005). A number of lines of evidence (Table 2) have been cited in support of an attack by a fungal disease (such as Ophiostoma) spread by the elm bark beetle (Scolytus scolytus L.). It is argued that the elm decline may be the first example of anthropogenic disturbance facilitating the spread of a pathogen (Lamb and Thompson, 2005). Evidence for such activity is often, but not always, associated with the major reduction in Ulmus pollen values and early agriculturists could

**Table 2.** Evidence offered in support of fungal diseases and/or insect infestations being contributory factors in the mid-Holocene declines of hemlock (*Tsuga*) and elm (*Ulmus*) pollen.

- 1. The apparent synchronicity of the events across broad geographical regions (see Davis, 1981; Huntley and Birks, 1983; Birks, 1986; Bennett and Fuller, 2002; Parker *et al.*, 2002).
- 2. The rapidity of the declines in *Tsuga* and *Ulmus* pollen values, as recorded at sites where high-precision dating evidence is available (e.g. Allison *et al.*, 1986; Peglar, 1993; Skog and Regnell, 1995).
- 3. The long-term impact on the abundance of hemlock and elm. At some sites *Tsuga* does not recover for 2 000 years (Davis, 1981) or in the case of *Ulmus* not at all (Birks, 1986). However, it should be noted that the late Holocene abundance of elm in Europe is likely to have been heavily influenced by human activity (Birks, 1986).
- 4. Fossil evidence for the contemporaneous presence of phytophagus insects. For the *Tsuga* decline, remains of the hemlock looper (*Lambdina fiscellaria* Gven.) and the spruce budworm (*Choristoneura fumiferana* Clem.), along with hemlock needle damage indicative of hemlock looper feeding, have been found coincident with the decline (Bhiry and Filion, 1996). The elm bark beetle (*Scolytus scolytus* L.) has been recorded before, and from around the time of, the *Ulmus* decline at sites in England, Scotland, Switzerland and Denmark (Girling, 1988; Rasmussen and Christensen, 1997; Clark and Edwards, 2004).
- 5. Modern outbreaks of virulent strains of fungal pathogens (e.g. chestnut blight, Dutch elm disease) have produced pollen stratigraphic signatures comparable to the mid-Holocene declines (e.g. Brugam, 1978; Davis, 1981; Perry and Moore, 1987).
- 6. The continued abundance of elm at a site in western Ireland through into the late-Holocene, despite evidence of localized disturbance due to human activity in the mid-Holocene. Lamb and Thompson (2005) suggest that the pathogen, vector or the tree was less susceptible near to the margins of the range of elm.

have been involved directly (by transporting a fungus) or indirectly (e.g. through causing damage favouring infection). Girling (1988) notes that although *S. scolytus* largely attacks dead wood, recently emerged beetles feed on healthy trees which can lead to their infection. Therefore, at some locations, for example where there is no association with human activity, disease might be a primary cause (Rasmussen, 2005). Alternatively, given that the elm decline is often accompanied by reductions in the abundance of other deciduous taxa (e.g. *Tilia*, Grant *et al.*, 2011) and the evidence, from a number of independent sources, for climate variability in Europe during the mid-Holocene (e.g. Langdon *et al.*, 2003; Magny *et al.*, 2006; Olsen *et al.*, 2010; Kühl and Moschen, 2012), the primary stress factor could have been climate change.

Recently obtained evidence relating to pathogen outbreaks during the Holocene include the investigations of Jasinski and Payette (2007) from sites in Québec. These authors have demonstrated the presence back to ca 8000 cal a BP of the spruce budworm (Choristoneura fumiferana Clem.) and the fungus Lophodermium piceae ([Fckl.] Hoehn.), which is associated with the needles of Picea mariana (Mill.). It is suggested that the co-occurrence of the budworm remains and fungal fruiting bodies indicate that spruce budworm infestations resulted in an increase in fungal infection and/or fructification, although drought is acknowledged as an alternative stress factor. In addition, remains of the spruce bark beetle Dendroctonus spp. have been reported from two lakes in the northern Rocky Mountains, USA (Brunelle et al., 2008). The evidence is compatible with infestations occurring ca 8 200 cal a BP, during a short-lived period of cooler and wetter conditions (e.g. Alley et al., 1997; Daley et al., 2011). This is consistent with the findings of Sherriff et al. (2011) who, using tree ring data for the period 1600-2007, showed that outbreaks of Dendroctonus rufipennis in southern Alaska were related to climate variability and, over multidecadal time scales, preceded by cool-phase Pacific Decadal Oscillation conditions.

Finally, fungal spores are increasingly being recorded alongside pollen in sedimentary records. High values of Kretzschmaria deusta ([Hoffm] P.M.D. Martin), a mild parasite of deciduous trees, have been reported from a number of sites. They occur after trees are likely to have been subject to stress from flooding (Kuhry, 1985; Waller and Grant, 2012) or human activity (van der Wiel, 1982) and have notably been recorded at a site in north Wales after the elm decline (Innes et al., 2006). In addition, Latałowa et al. (2013) report the sustained abundance of K. deusta spores over the interval ca 8 000-5 300 cal a BP from a mire in northern Poland and suggest that the perpetual activity of this fungus may have influenced vegetation composition. High frequencies of Tilia pollen (and therefore warm summers) and coprophilous fungal spores are also recorded during this period, with drier conditions and animal damage likely to have contributed to the infection of Tilia by K. deusta. The ability of Tilia to respond to these attacks by the vegetative production of basal sprouts could not only have enabled the long-term survival, but also promoted the local dominance, of Tilia (Latałowa et al., 2013).

## Towards a better understanding of the long-term history of disturbance events

The investigations undertaken to date suggest that palaeoecology can contribute to a better understanding of forest disturbance episodes involving pathogens, both large-scale and gap-phase events. However, this potential has yet to be fully realized. Pathogen-focused research strategies, informed by an understanding of the interactions and chain of events within which pathogens are active, are required. In conjunction with multi-site pollen investigations undertaken at an appropriate temporal resolution, additional complementary evidence needs to be collected (Table 1).

Wood remains have considerable potential. Tree ring analysis offers a direct means of demonstrating stress. For example, Bhiry and Filion (1996) report decreases in radial growth associated with recent hemlock lopper outbreaks in Québec and in fossil *Tsuga* remains dating back to ca 5 650 cal a BP. Periods of accelerated growth due to the death of mature trees and release from competition have also been identified after recent and historical spruce bark beetle outbreaks in Colorado and Alaska (Veblen *et al.*, 1994; Sherriff *et al.*, 2011). Wood remains may also record the pattern of damage caused by bark beetles and fungal attack. For example, Creber and Ash (1990), on the basis of welldefined pockets of decay, were able to attribute damage on Upper Triassic trees in the south-western USA to a pathogenic pocket rot fungus rather than a saprophyte.

Due to the growth in the use of non-pollen palynomorphs for palaeoenvironmental reconstruction and advances in fungal spore identification (e.g. van Geel, 1978, 2001; van Geel and Aptroot, 2006), as previously noted, fungi are increasingly being recorded in association with forest disturbance events. A number of taphonomic issues need to be considered. Fossil records will clearly be biased towards fungi aerially transmitting robust and distinct spores with diseases transmitted by insect-vectors likely to go unrecorded. The spores of Ophiostoma are, for example, fragile and unlikely to withstand the procedures used to prepare fossil pollen and spores (Groenman-van Waateringe, 1983). In addition, van Geel and Andersen (1988) have demonstrated that the frequency of K. deusta spores, which are produced close to the ground, is influenced by the distance between sampled sites and infected trees and Waller (2010) presents evidence indicating that fungal spore representation can be strongly influenced by changes in depositional environment. It should also be noted that the distinction between pathogens and saprophytes in fungi can be problematic. As noted earlier for Armillaria, many fungi, while they may act as parasites, continue to, or mainly, exploit dead hosts.

For the elm decline, particular caution needs to be taken over inferring the presence of a fungal disease from finds of the elm bark beetle (Table 2). Not only are the beetle remains often weakly or not associated with the decline in pollen values, but with the beetle present earlier in the Holocene, fossil occurrence may simply indicate the presence of suitable habitat (Robinson, 2000).

Whether climate change acted as a primary causal agent of a disturbance episode is a testable hypothesis if independent palaeoclimatic data can be obtained (cf. Zhao *et al.*, 2010). Comparison of high-resolution pollen and stable isotope data could, for example, confirm or refute a climatic trigger for the elm decline.

Pathogen attacks are likely to result in an increase in combustible material and therefore to be associated with fires, a connection which can be examined through investigations of charcoal content. Jasinski and Payette (2007) found *L. piceae* remains, along with spruce budworm head capsules, before fire events. In addition, in a well-resolved sequence from Brede Bridge in southern England (Grant and Waller, 2010), a peak in micro-charcoal frequencies occurs approximately 100 years after the start of the elm decline. Such a figure is consistent with an event involving the protracted decline and death of trees.

#### The importance of pathogens in the past

From the evidence currently available it could be concluded (cf. Davis, 1981) that large-scale pathogen outbreaks were much more infrequent in the past which, given the role that human activity currently plays in transporting pathogens to new locations, would not be surprising. Multi-site, highresolution, multi-proxy palaeoecological investigations require considerable resources. Can such studies into past pathogen outbreaks be justified? Causal connections between biological trends and climate change are often difficult to establish from short-term studies, and palaeoecology has a role to play in demonstrating the degree to which climate change and pathogen attacks are coupled. The evidence currently available certainly hints at a link with abrupt climate change, although associations between the various proxies are often weak.

Pathogens, even if demonstrated to have been active during disturbance events in the past, could be regarded as simply being secondary factors and their appearance the inevitable consequence of the presence of stressed trees. However, even if not the primary stress factor, pathogens may still have significantly influenced vegetation composition. One of the characteristics of the hemlock and elm declines is that the reductions in the abundance of Tsuga and Ulmus pollen are long-lived (Table 2). In contrast, associated taxa often decline simultaneously but then recover. Given that pathogens tend to be species specific, by contributing to tree death and the failure of Tsuga and Ulmus to regenerate, they may in some areas have been the critical factor in causing their hosts to disappear. Host trees in less stressed positions would be vulnerable because of the increased loading of pathogens, with re-infection preventing re-establishment through vegetative means or propagules. Recovery would then have been delayed until resistant stock could reinvade from distance or the virulence of the pathogen was reduced (Table 1). Multiproximity studies have the potential to address this issue; is the long-term impact of climate change on vegetation composition greater when accompanied by a secondary pathogen attack?

In part, the hemlock and elm declines stand out because of their geographical scale. However as Birks (1986) points out, Holocene pollen records from Europe and North America document many other smaller-scale (regional and local) reductions in the abundance (or extinctions) of tree taxa. Given the role played by non-exotic, as well as introduced, pathogens in influencing the modern distribution of species (Castello et al., 1995; Holdenrieder et al., 2004), multi-proxy investigations are likely to demonstrate that pathogens were an important factor in at least some past changes in forest composition and ultimately to contribute to a better understanding of such episodes. Such knowledge is clearly important as, whether driven directly by human activity or by climate change, ash dieback seems destined to be but one of an increasing number of high-impact pathogen outbreaks.

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