**Working title: Positive feedbacks cause collapse of tree cover in a temperate forest**

Main message of paper: Transition to non-forest state depends on positive feedbacks: loss of mature trees causes lower recruitment and tree death increases likelihood of death of nearby trees

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**Summary**

1. There is concern that forest dieback may lead to transitions to non-forest states. Positive feedbacks are thought to play an important role in such transitions by creating self-perpetuating shifts in system states. Despite this concern, there is has been relatively little work on the topic.
2. We used statistical models to identify correlates of death and recruitment of the canopy dominant (*Fagus sylvatica* - beech) in a temperate forest site that has been sampled over 50 years and appears to be undergoing transition to grassland. We use these results along with information from the literature to build an individual based model to investigate the impact of positive feedbacks on forest persistence.
3. We found that the probability of tree death declined with increasing tree growth rate, distance to nearest dead tree and sand content of soils, but was positively correlated with tree size. Seedling density was negatively related to canopy cover, but sapling density was higher in areas of high canopy cover.
4. Contrary to our expectations, inclusion of potential feedbacks in our individual based model did not result in total loss of tree cover over 150 years, even with 100% mortality of juvenile trees.
5. Our results suggest that while positive feedbacks may influence both recruitment of juveniles and death of mature trees, the long lifespan of trees provides a buffer to sudden transitions. However, under current conditions the woodland may transition to a savannah-like system.
6. **Synthesis and applications:** To enhance forest resilience management should attempt to stop the development of such positive feedback loops. In our study area fencing off forest areas to reduce seedling mortality caused by browsing of ponies and deer may reduce feedbacks related to juvenile death. However, reducing the feedbacks related to mortality of larger trees will be more challenging as these may be related to larger scale drivers relating to climate change.

**Introduction (aim for ~800 words)**

There is widespread concern that forest dieback may result in shifts to treeless states (Reyer *et al.* 2015). Over the past decade research has suggested such shifts may occur in both tropical (Barlow & Peres 2008; Hirota *et al.* 2011) and boreal regions (Scheffer *et al.* 2012) as a result of a both changes in climate and disturbance regimes. Forests may be particularly vulnerable to rapid changes because trees are long lived, immobile organisms, that consequently find it difficult to adapt to new environmental conditions (Burrows *et al.* 2011; Seidl *et al.* 2015). Any shift to relatively treeless, non-forest states would cause loss of forest biodiversity as well dramatic changes in the ecosystem services provision (Scholes *et al.* 2014). Due to these risks a recent IPCC assessment concluded that forest dieback has the potential to cause major economic impacts (Scholes *et al.* 2014). However, despite these concerns there is relatively little known about the mechanisms that cause transition of forests to relatively treeless states (Reyer *et al.* 2015).

Transitions from one ecosystem state to another occur when disturbances (*sensu* Sousa 2001) resulting in death of organisms cause changes to a system from which it fails to recover (Nimmo *et al.* 2015). These disturbances may represent instantaneous or continuous alteration of species abundances, which are respectively termed pulse and press disturbances (Bender, Case & Gilpin 1984). Critically, what is considered to be ‘instantaneous’ or ‘continuous’ depends on the generation time of the organisms in an assembly (Shade *et al.* 2012). Theory relating to ecosystem resilience suggests that interactions between different drivers may result in positive feedbacks driving the system into a different state as a result of rapid non-linear regime shifts (Scheffer *et al.* 2001). However, linear, gradual changes in ecosystems resulting in a shift to a novel state also occur (Davidson 2000; Petraitis & Hoffman 2010; Petraitis 2013), though have received less attention in the ecological literature. Importantly, the presence of positive feedbacks may make recovery of a system back to a desired state difficult, while linear change may be more easily reversed via management (Folke *et al.* 2010).

From the perspective of forest conservation, feedbacks of particular concern are those between local disturbances (e.g. fire, pests, drought or deforestation) and climatic changes that impair regeneration of tree species (Reyer *et al.* 2015). For example, logging and deforestation in tropical forests combined with drought and increased fire frequency may lead to a shift to a savannah-like vegetation structure (Nepstad *et al.* 1999; Barlow & Peres 2008). Similarly, large disturbances in Mediterranean forests can lead to reduced seedling recruitment and invasion by grasses and shrubs, which result in increased fire frequency and further suppression of tree cover (Acácio *et al.* 2007). However, dieback is also occurring in temperate forests that do not exhibit similar feedback mechanisms (Martin *et al.* 2015). Tree mortality as a result of climate change, pathogens and insect pests is increasing in many temperate forests (van Mantgem *et al.* 2009; Seidl *et al.* 2014) and so identification of potential feedbacks that may drive these systems into non-forest states is critical to inform management decisions.

Here we use a long-term data set, collected in a temperate forest ecosystem that has undergone partial stand dieback in recent decades. At this site in southern England, monitoring data have been collected repeatedly over a period of 50 years from 1964-2014 (Mountford *et al.* 1999; Mountford & Peterken 2003; Martin *et al.* 2015). Using statistical models we investigate correlates of tree mortality and recruitment, both of which include potential feedback mechanisms. We then use results from these models and information from the literature to produce an individual based model to test the effects of identified feedbacks on forest structure. Specifically, using these methods our aims are:

1. To test the factors that influenced seedling and sapling density
2. To test the influence of tree size, growth rate, proximity of dead trees and soil characteristics on mature tree mortality
3. Test the influence of feedbacks identified in statistical models on forest structure using an individual based model

**Methods (aim for 1500 words, currently 2120)**

**Site description**

For this study we used data collected over a 50 year period in Denny Wood, which is located in the New Forest National Park, in Southern England (Lat: 50o 51.5’ N and 1o 32.5’ W). The site is gently sloping (1-3 degrees), with clay rich brown earth soils. Woodland vegetation is dominated by old-growth beech (*Fagus sylvatica*) with frequent pedunculate oak (*Quercus* robur) and birch (*Betula pendula, B. pubescens*), and an understory primarily of holly (*Ilex aquifolium*). In open areas the ground vegetation is mostly comprised of *Agrostis*-dominated grassland or stands of bracken (*Pteridium aquilinum*). There are large populations of deer, ponies and cattle in the New Forest, resulting in high herbivore pressure. The site which we used for this study is likely to have experienced high browsing pressure since the 1960’s (Mountford & Peterken 2003). More detail on the site is given in Mountford *et al* (1999), Mountford and Peterken (2003) and Martin *et al* (Martin *et al.* 2015).

**Data collection**

Measurements were conducted in one 20 m-wide 1 km long transect, which was originally established in the 1950s. The transect was subdivided into contiguous 45 20 x 20 m (0.04 ha) subplots and surveyed in 1964, 1984, 1988, 1996 and 2014. Details of measurements are presented Mountford *et al* (1999), Mountford and Peterken (2003) and Martin *et al* (Martin *et al.* 2015). In each survey, the location and species name of all woody stems >1.3 m in height were recorded, their diameter at breast height measured using diameter tapes, and their status assessed as either alive or dead. Each stem >1.3m height was given a unique ID number to allow individual trees to be tracked. Stems <10 cm DBH were classified as saplings and those >10 cm DBH as mature.

In 2014 we collected data on seedling density, canopy openness, and soil characteristics. The density of tree seedlings of all species present in 10 x 10 m plots located in the centre of the 20 x 20 m plots was recorded. Canopy openness of subplots was assessed using a concave spherical densitometer in all four corners and the centre of 20 x 20 m plots, and the mean calculated for each subplot. Soil type was assessed by collecting 3 soil samples from within each 20 x 20m subplot using a 5 cm diameter soil corer. The first 20 cm of the mineral layer was retained. Soil samples were sent to the Forest Research laboratories in Surrey, UK where particle size distribution of the soil was determined by suspending 30 g of soil in water which was passed through the flow cell of a laser diffraction particle size analyser (Beckman Coulter LS13320).

**Statistical analysis**

**Recruitment of juveniles**

Our assessment of tree recruitment focussed on saplings (woody stems >1.3 m height and <10 cm DBH) and seedlings. We tested for relationships seedling density and canopy openness and grazing pressure using generalised linear models. We also tested the the effect of the density and basal area of mature trees (stems >10 cm DBH) on sapling density using generalised mixed models with a poisson error structure for the three major woody species present: beech, holly and oak. These models included interactions with survey year to identify if the slope of the relationship between mature tree density and sapling density changed over time. To determine whether the widespread loss of smaller stems seen in a previous study at the site (Martin *et al.* 2015) was largely attributable to growth of individuals into other size classes or mortality we tracked the fate of each sapling recorded. As part of this we also calculated the annual mortality rate, , as defined by Sheil, Burslem & Alder (1995):

where and are the number of stems at the first and second surveys respectively and is the number of years between censuses.

**Individual tree mortality**

We assessed whether some of the mature tree mortality observed in Denny wood could be explained by the self-thinning process using a linear mixed model with subplot number as a random effect to relate stem density to basal area. A negative slope suggests a gain in BA with a loss of stem density while a positive slope suggests a gain in BA with increasing stem density.

During data processing tree status (alive or dead) was coded at census time. Only trees with diameter measurements at both and prior to time were included in analyses. Censuses on the transect were undertaken 5 times from 1964-2014 with a mean (± SD) census interval period of 12.5 ± 6.7 years (range 4-20 years). Since trees used in mortality models required censuses at both and prior to time we selected three non-overlapping census periods: 1984-1988, 1988-1996 and 1996-2014 (mean census period 10 ± 5.9 years). Statistical models of individual tree mortality were developed using logistic mixed effects models, which describe the probability that a tree dies in a given period of time. To correct for the variation in census interval we used a complementary log-log link with an offset equal to the census interval so that predictions from models were equivalent to the annual probability of mortality (Fortin *et al.* 2008). Subplot ID number was used as a random effect to account for repeated sampling of the same plots (Fortin *et al.* 2008).

Models were developed in a four-step process similar to the workflow of Chao et al. (2008). In step 1 we prepared predictors classified into 4 groups:

* 1. tree size – DBH (cm), Basal area (m2), and tree size relative to other trees in the transect (bounded between 0 and 1)
  2. tree growth - Annual diameter growth rate (mm year-1), basal area growth rate (mm2 year-1), relative DBH growth rate (% DBH increase year-1), and relative BA growth rate (% BA increase year-1)
  3. proximity to dead trees - distance of an individual tree to a dead tree and abundance of dead trees in a 10m buffer
  4. soil type - percentage of each soil sample classified as sand

For more detail on the calculation of these variables see the supplementary materials. All model variables were standardised using the methods of Schielzeth (2010) by subtracting the mean of the variable and dividing by its standard deviation. This allows coefficients to be interpreted as effect sizes, reduces collinearity between variables and improves model convergence (Schielzeth 2010).

In step 2 we selected the best predictor for each group by choosing the univariate logistic mixed effect models which had the lowest AICc (Burnham & Anderson 2002; Chao *et al.* 2008a). This step reduces intercorrelation of variables which can lead to difficulty in interpreting effects (Chao *et al.* 2008b). In step 3 a full multivariate model was developed using these selected variables using additive terms only. In step 4 model averaging was used to produce parameter estimates for models with a ΔAICc≤7.

**Individual based model**

We used an individual based model, built using Netlogo (Wilensky 1999) to investigate the importance of feedbacks in causing collapse of forest structure. The model description follows the ODD protocol for describing individual based models (Grimm *et al.* 2006).

**Purpose**

We used the model to investigate under what conditions loss of tree cover and basal area (BA) might occur in a simplified representation of a New Forest beech woodland. The only species represented is beech, as this is the dominant species found in the area, and mortality of the species has caused the majority of BA loss from 1964-2014 (Martin *et al.* 2015).

**Entities, state variables and scales**

The model comprises of two types of entities: grid cells and individuals. Individuals represent beech trees. Each individual is characterised by its location, development stage (juvenile or mature), age (in years), DBH (cm), basal area (m2), mean seed dispersal distance (mean distance from the source, in number of cells), growth rate in previous year (mm year-1), local basal area and stem density of mature trees (within a circles with an area of 400 m2, the size of plots we used for surveys), and distance to nearest dead tree (m) . DBH of mature trees and juveniles is derived from the age of trees using an equation for beech growth defined in Holzwarth *et al.* (2013) and BA defined as . Local basal area and stem density represent the total BA and stem density within a circle with an area of 400 m2 – equivalent to the plot size used in data collection. Dispersal distance is a random number drawn from an exponential distribution with a mean of 5 m.

All grid cells in the model are considered suitable for individuals. Each grid cell is characterised by its location, whether a tree has died in that patch and the time since last tree death on that patch. When a tree dies the patch value changes from 0 to 1 and after 10 ticks if no other tree has died on this cell this value returns to 0. Each grid cell can only contain one mature individual, but may contain multiple juveniles. The model landscape consists of 100 x 100 grid cells, with each cell representing 1 m2, thus the entire area represents 1 ha. Each model time step represents one year.

**Process overview and scheduling**

Initially the distribution of individuals is determined by randomly distributing 280 mature individuals with a random age drawn from an exponential distribution with a mean of 80 years assigned to each individual. This was approximately the density and age structure of Denny Wood when first surveyed in 1964. At the same time 100 juvenile trees are randomly distributed across the space. Then in each time step the following events are processed in the given order: increase age of individuals by one year, increase individual DBH & BA, identification of whether the time step represents a mast year, seed dispersal from mature trees > 40 years old, and death.

**Design concepts**

The total number of mature trees and basal area *emerge* from changes in the probability of mature tree mortality that occur as they age and increase in size, as well as from changes in the mortality of juvenile trees. *Interactions* between individuals are the result of density dependant mortality processes, which show size asymmetry. For juveniles this is modelled by defining a maximum number of juveniles that can coexist in the area as 1000, and a maximum number of juveniles per cell as 100. When these numbers are exceeded the smallest juveniles are killed. Similarly, for mature trees the local maximum BA was set at 75 m2 ha-1, the maximum observed for any plot during 1964-2014. When this maximum is exceeded the smallest mature tree in an area of 400 m2 is killed.

In addition our model tests the impact of reduced survival of juveniles in areas with few mature trees, as seen at our site, by allowing the user to switch this on and off. When turned on this causes juveniles in locations where mature tree density is <5 in the surrounding 400 m2 to have a 0.02 annual chance of survival, based on our observations that no saplings (trees >1.3 m tall, <10 cm DBH) were found in areas with tree density lower than this. Additionally, our model tests what impact an increase in the probability of mature mortality as a result of being close to a dead tree would have on forest structure. This can be switched on and off. When switched on this causes the annual mortality to be increased as a function of distance to nearest cell with a dead tree. The parameter estimate for this was derived from our statistical model of mature tree mortality described above.

*Stochasticity* is used in the model to define whether a given time step represents a mast year. Beech trees produce large amounts of seed once every 2-3 years in the UK (Packham *et al.* 2012) and thus we set a probability of 0.3 of each year being a mast year.

To test the model we first initiated the model so that forest structure was similar to that seen in 1964. We then ran different scenarios with and without differential juvenile mortality and spatial mortality feedbacks and compared the change in BA and forest structure after 50 time steps to those we observed in 2014. Following this we ran the models using the same parameterisation for a further 100 years to investigate possible future changes in BA and tree cover under the different scenarios. Each model run had 100 iterations and median values were used to summarise model results.

**Results (aim for ~600 words) – currently 539**

**Tree recruitment**

Mean beech seedling density (± SE) in 2014 was 115.22 ± 32.14 seedlings ha-1. Canopy openness was positively related to beech seedling density (slope=0.56 ± 0.09, P value <0.001), but no other variables were included in models which had ΔAICc≤7 and thus were considered to have poor support. No metrics of deer or pony density were related to seedling density in any way.

There was a negative relationship between sapling density and canopy openness. Only 6 plots had any saplings present, but all of these plots had a canopy openness of < 20%. There was a positive relationship between mature beech density and sapling density (slope= 1.27 ± 0.14, P value<0.0001), and sapling density tended to be reduced over time (slope= -0.69 ± 0.10, P value<0.0001). The relationship between mature beech density and sapling density grew more positive over time (interaction term=0.38 ± 0.09, P value<0.001), though since the density of mature trees was also reduced over this time period this effect was relatively unimportant in determining sapling densities. As the number of beech saplings declined during the years 1964-2014, so did the mortality rates of these saplings, from a maximum of 4.07% per year in 1964-1984 to 0.50% in 1996-2014 (Table 1). Conversely the proportion of saplings that became mature trees (>10 cm DBH) showed an increase over this time period (Table 1).

**Tree mortality**

The slope of the relationship between log subplot stem density and log subplot basal area was positive (slope=0.41 ± 0.05, marginal R2=0.24, Figure 1). However, in general subplots lost both stem density and basal area between 1964 and 2014 (Figure 2). Given that self-thinning processes tend to be strongest when plots are increasing in biomass and losing stem density at the same time (Coomes & Allen 2007) such processes are unlikely to be responsible for the majority of tree death seen in Denny wood from 1964-2014.

When predicting the mortality of individual beech trees growth rate was considered the most important predictor, as it was included in all models with a ΔAICc≤7. Trees that grew slowly or shrunk were more likely to die than those that grew relatively quickly (slope=-0.59 ± 0.06, P value <0.001, Figure 3a). Next most important was tree DBH with an importance value of 0.8, and models suggested that tree size was positively correlated with probability of mortality (slope=0.21 ± 0.05, P value<0.001, Figure 3b). Distance to nearest dead tree and soil type were of similar importance with importance values of 0.52 and 0.45 respectively, with models indicating that trees closer to dead trees were more likely to subsequently die (slope=-0.24 ± 0.06, P value<0.001, Figure 3c) and trees located in areas of the forest where soils had higher sand content were less likely to die (slope=-0.27 ± 0.01, P value<0.001, Figure 3d).

**Individual based model**

Results from our individual based model suggest that when the annual probability of juvenile death is low, the forest does not undergo a transition to a treeless state even if feedbacks are present. However, when annual probability of juvenile death is >0.4 these feedbacks are enough to push the system into a decline, particularly when both distance to nearest dead tree causes an increase in the probability of mortality and juvenile survival is reduced in gaps (Figure 4). Thus the effect of feedbacks was dependent upon high background juvenile mortality.

**Discussion (aim for ~1200 words)**

1. Summary of results

In our study we found that recruitment of beech seedlings and saplings in Denny wood was limited. Seedling density was negatively correlated with canopy cover, while sapling abundance was positively correlated. Modelling of mortality showed that slow growing, large trees, close to dead trees, on soils with low sand content were the most likely to die. An individual based model suggested that forest basal area would decline unless juvenile mortality was dramatically reduced. However, none of the scenarios suggested the potential for total loss of tree cover.

1. Causes of juvenile recruitment limitation

Seedling density was highest in areas with low canopy cover reflecting increased germination and growth rates of tree seedlings where light intensity is higher and grass or shrubs provide protection against ungulate herbivores (Collet, Lantera & Pardos 2001; Kuijper *et al.* 2010; Duwyn & MacDougall 2015). However, this was not reflected by higher sapling density in areas with low canopy cover, but the opposite resulting in very limited regeneration following death of large trees (Martin *et al.* 2015). There are a number of potential explanations of this. It is possible that ungulate browsing is concentrated in areas where grass cover is highest resulting in high mortality of seedlings which are eaten. Meanwhile, seedlings regenerating under closed canopies may be less likely to be eaten. Though grass can provide protection from browsing herbivores, at high densities this protection can be overwhelmed (Duwyn & MacDougall 2015). Given the high density of herbivores in the New Forest, this appears to be the most likely cause of recruitment failure in the woodland we used in our study. However, debarking of beech saplings by grey squirrels has also been seen in Denny wood, suggesting a further mechanism limiting recruitment.

1. Causes of death

The strongest predictor of tree death was its growth rate, with slow growing trees more likely to die. Previous work has shown that beech trees that grow slowly are more likely to die (Holzwarth *et al.* 2013), and this correlation may reflect increased mortality under unfavourable growing conditions as well as senescence. The second most important predictor of beech mortality was DBH. This positive relationship between DBH and mortality indicates a stand undergoing disturbance, in our case probably as a result of the combined effects of summer drought, winter waterlogging, disease and overgrazing (Martin *et al.* 2015). Interestingly, tree proximity to dead trees was positively related with tree mortality. The mechanism behind this relationship is unclear but may relate to increased stress on trees close to gaps where no regeneration has occurred, similar to edge effects. It is also possible that fungal pathogens such as *Phytophthora*, which has caused beech mortality in continental Europe and has been recorded in the New Forest, may result in this mechanism by spreading amongst trees. The percentage sand content of soils was negatively related to beech mortality, suggesting that winter water-logging and summer drought may have been partially responsible for death of mature beech trees.

1. Feedbacks may be important in our system, as seen in other transitions (~200 words)

Our analyses indicate that positive feedbacks are potentially important for structuring the forest system. Our observations that seedling density was highest in areas with low canopy cover but sapling density was highest in areas with high canopy cover suggests such a feedback. This observation suggests that mortality mechanisms for seedlings and saplings may differ resulting in clusters of treeless areas. This has the potential to result in a much more open woodland than that found at present, as suggested by Vera et al. The mechanism for this may be that the high ungulate density in the New Forest causes total death of seedlings in gaps, where ponies and deer feed preferentially. In addition, proximity to dead trees resulted in an increased probability of mature tree death. If tree recruitment is higher under areas of high canopy cover this could ultimately result in an increase in stand level mortality.

1. However, rapid collapse to a treeless state does not occur because trees buffer against this (~200 words)

Despite the potential importance of feedback loops in driving change in our site, simulations of the next century never resulted in a transition to a completely treeless state. In addition, declines in forest basal area were largely linear and did not follow the trajectory associated with regime shifts of a sudden decline. This suggests that while the forest may be undergoing dieback the current level of disturbance is not high enough to result in a loss of tree cover in the foreseeable future. However, disturbance intensity from climate and pests are predicted to increase in European forests in the coming century (Seidl *et al.* 2014), meaning predictions from our model may be conservative. This lack of loss of forest cover in our study was a result of the long generation times of trees, which helped to buffer against total loss as a result of disturbance. This is synonymous with extinction debt where there is a time-lag between disturbance and the ultimate loss of a population. Such declines are likely to relatively common in forest, compared to mass mortality over relatively short period of time. In the context of forests, rapid loss of tree cover seems to require interactions between disturbance and fire. Such interactions have been observed in tropical (Barlow & Peres 2008), Mediterranean (Acácio *et al.* 2007), temperate (Flory *et al.* 2015) and boreal forests (Scheffer *et al.* 2012) suggesting a universal mechanism for feedbacks resulting in forest loss.

1. Managers should attempt to break these feedback loops or stop them from being established (~200 words)

**Conclusion (200 words)**

1. Feedback loops can be important in causing transitions to non-forest states and need to be managed carefully (~200 words)

Over the next 100 years, without appropriate management there is the potential for widespread loss of tree cover in areas of the New Forest undergoing dieback. Our estimates of 50-70% losses in basal area may be worsened by projected increases in temperature, drought frequency, and outbreaks of pests and diseases. Such die-off is occurring elsewhere in the New Forest and presents the risk of widespread losses of biodiversity and ecosystem services. The most obvious way to reduce the probability of die-off is to increase beech recruitment by protecting selected woodlands from overgrazing. The potential impacts of fungal pathogens and drought may also be reduced by this recruitment, resulting in a forest composed of trees with a wider range of tree ages.

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**Figures**

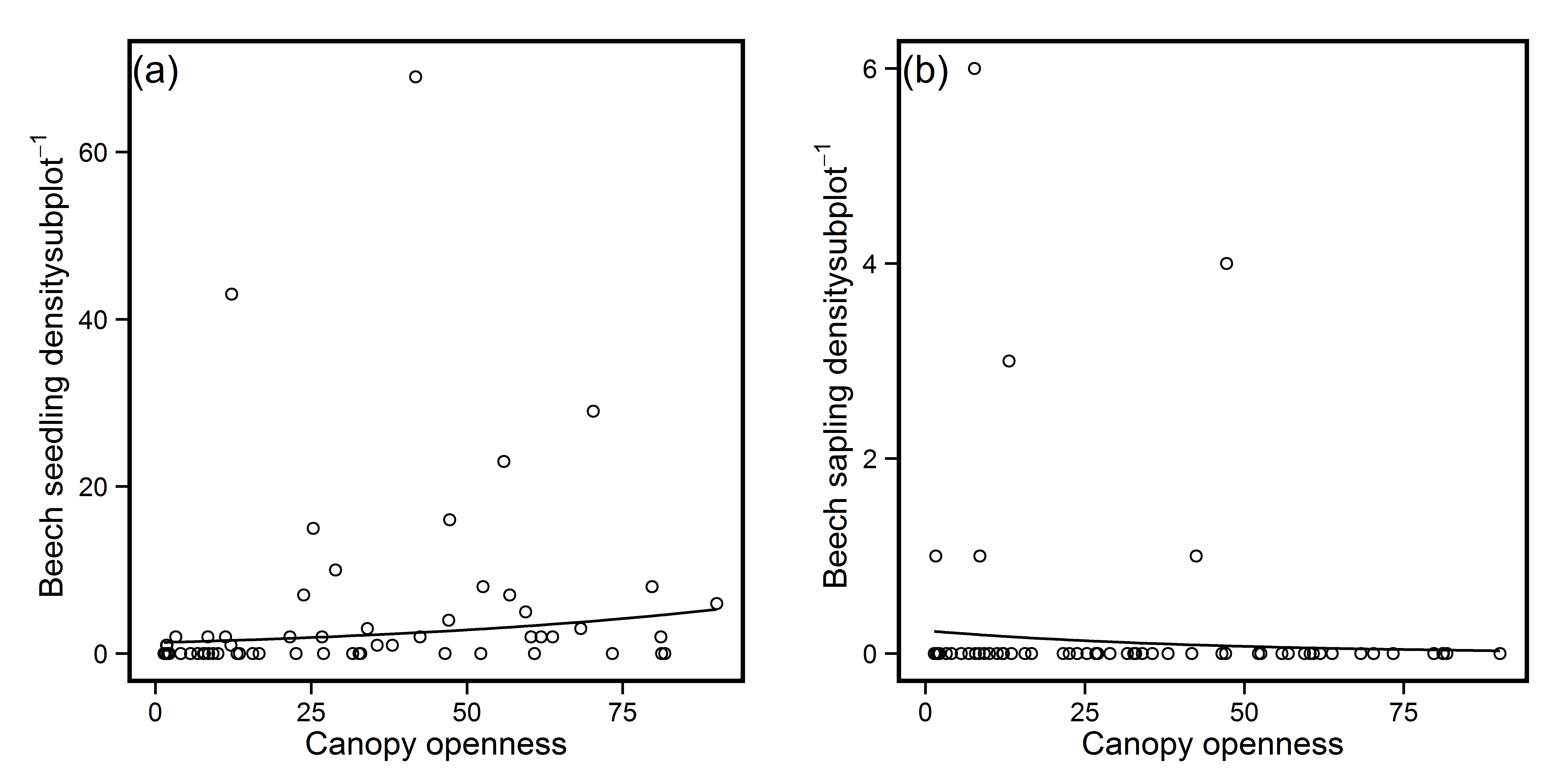


Figure 1 – Relationship between density of beech (a) seedlings and (b) saplings and canopy openness in woodlands in the New Forest showing signs of die-off. Solid lines represent predictions from averaging of mixed effects model coefficients.



Figure 2 – Relationship between subplot stem density and total subplot basal area. Points represent individual plots in 1964 (red circles), 1996 (green triangles) and 2014 (blue squares). The solid line represents the prediction from a mixed model of this relationship with the grey band representing the coefficient confidence intervals. Note that that both x and y axes are log transformed.

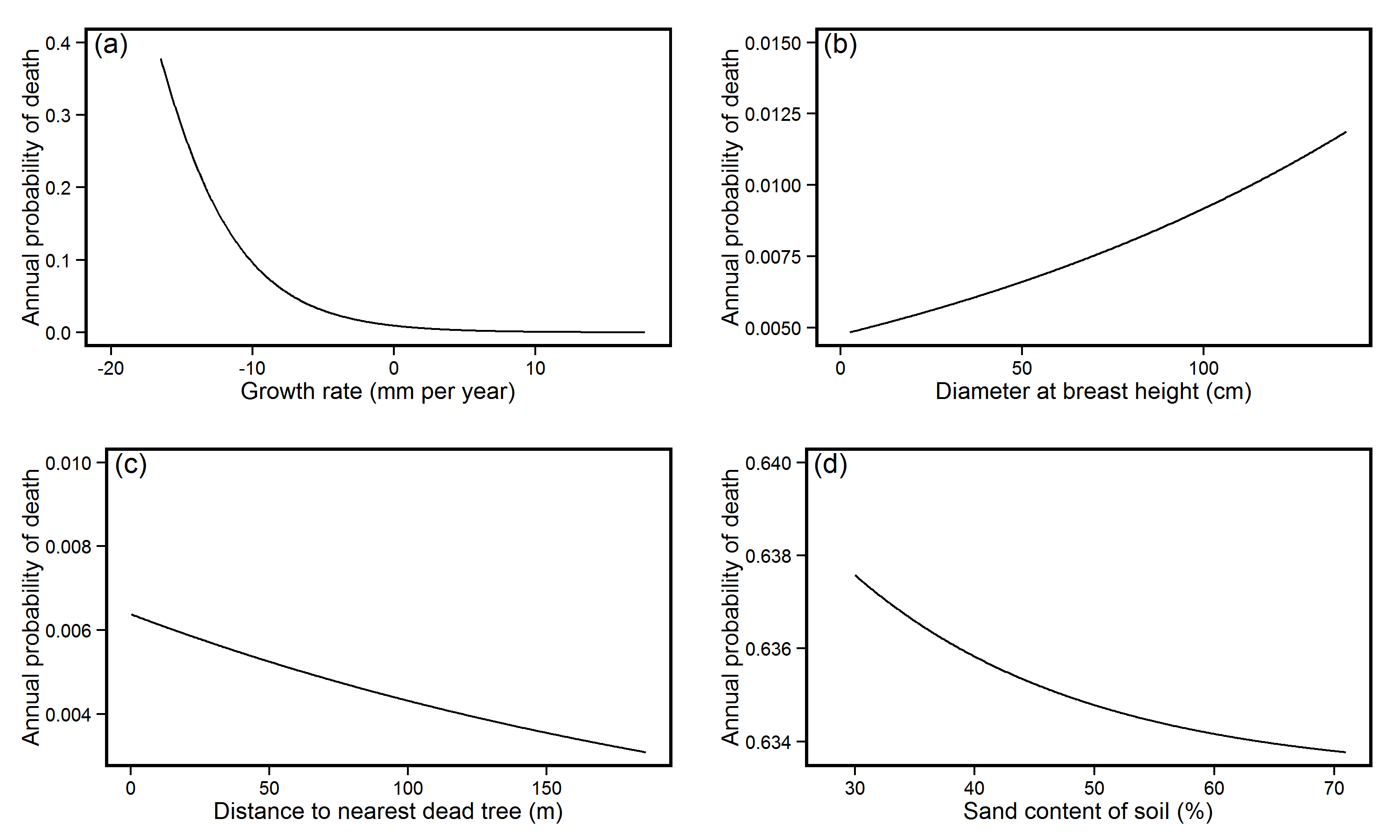


Figure 3 – Relationship between annual probability of beech tree death (a) growth rate per year, (b) diameter at breast height, (c) distance to nearest dead tree and (d) sand content of soil. Lines represent predictions generated from model-averaged parameter estimates.

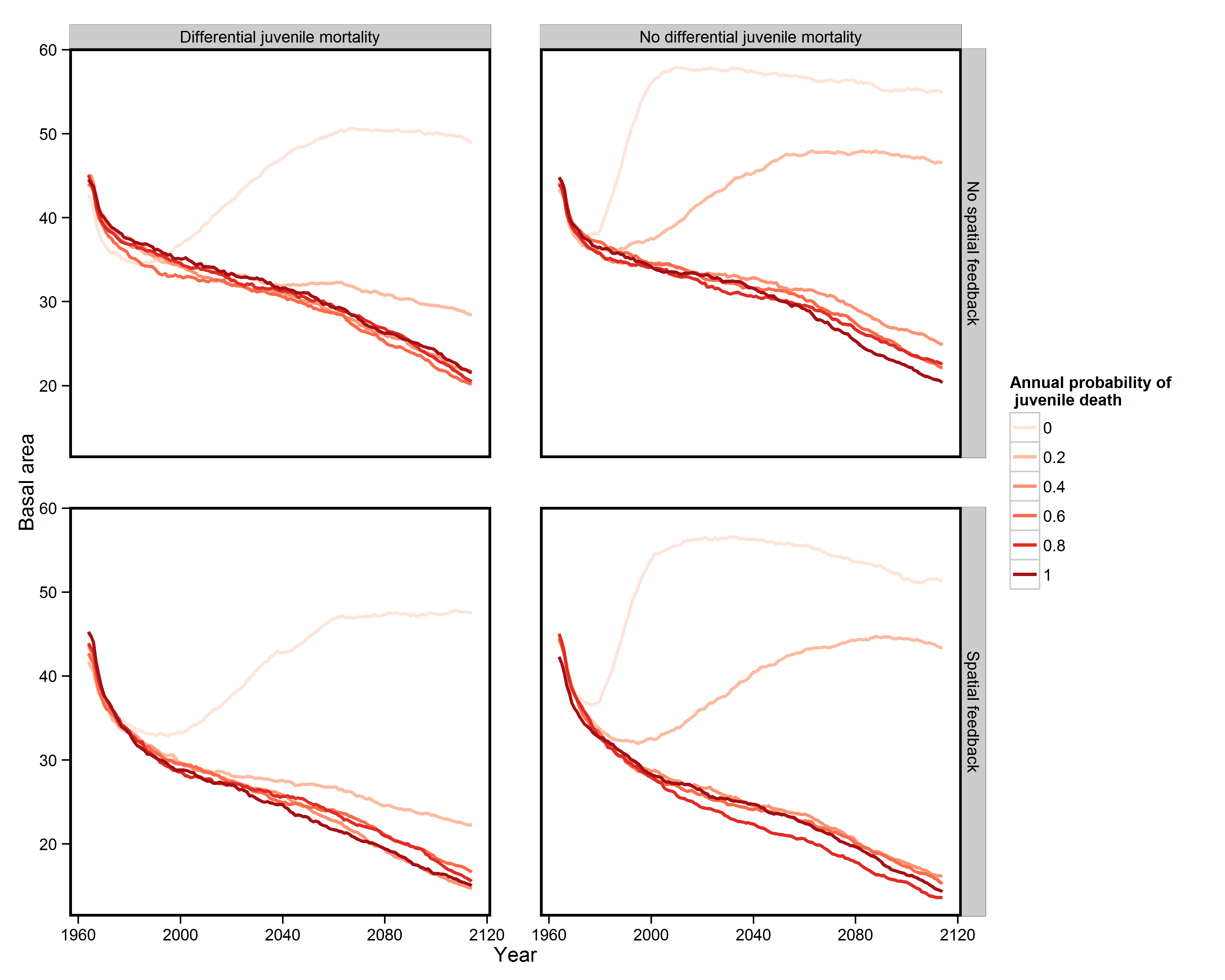


Figure 4 – The effect of feedbacks in mature tree death and juvenile mortality with varying probability of juvenile death on basal area from 1964 to 2114. Coloured lines represent mean basal area at each model time step with band representing the mean ± 1 SD. Black lines represent field observations from 1964 – 2014 to allow comparison between model results and real data.

Important that we show that even under high mortality you don’t get non-linearity

Table 1 – Summary of recruitment and mortality of saplings in Denny wood from 1964 to 2014.

|  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Census period | Proportion of saplings that died | | | Proportion of saplings that increased to >10 cm DBH | | | Annual mortality rate of saplings | | |
|  | Beech | Oak | Holly | Beech | Oak | Holly | Beech | Oak | Holly |
| 1964-1984 | 0.56 | 0 | 0.78 | 0.14 | 1.00 | 0.04 | 4.07% | 0 | 7.20% |
| 1984-1988 | 0.11 | NA | 0.22 | 0.2 | NA | 0.01 | 2.79% | NA | 5.91% |
| 1988-1996 | 0.15 | NA | 0.25 | 0.32 | NA | 0.08 | 2.01% | NA | 3.53% |
| 1996-2014 | 0.09 | NA | 0.62 | 0.61 | NA | 0.16 | 0.50% | NA | 5.22% |

Table 2 – Coefficients of beech tree mortality from 1964 to 2014 produced from model averaging of mixed effect complementary log-log models with ΔAICc≤7.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Variable** | **Estimate** | **Std. Error** | **Lower CI** | **Upper CI** | **p value** | **Importance value** |
| Intercept | -5.12 | 0.16 | -5.43 | -4.82 | <0.0001 | 1 |
| DBH | 0.21 | 0.05 | 0.11 | 0.31 | <0.0001 | 0.8 |
| Distance to dead tree | -0.24 | 0.06 | -0.36 | -0.12 | <0.0001 | 0.53 |
| Growth rate | -0.59 | 0.06 | -0.70 | -0.47 | <0.0001 | 1 |
| Sand content | -0.27 | 0.01 | -0.29 | -0.26 | <0.0001 | 0.45 |

Table 3 – Parameter values for individual based model

|  |  |  |  |
| --- | --- | --- | --- |
| **Parameter name** | **Sources** | **How derived** | **Value** |
| Number of seedlings produced in mast year per tree | (Olesen & Madsen 2008) | Mean of number of seedlings present after a mast year divided by the number of mature beech trees in the woodland. | 82 (26) |
| Juvenile height growth rate in gaps | (Collet *et al.* 2001) | Used values from reference | 9.5 cm year-1 |
| Juvenile height growth rate under closed canopy | (Collet *et al.* 2001) | Used values from reference | 1.2 cm year-1 |
| Maximum juvenile density | (Olesen & Madsen 2008) | Used values from fenced, ungrazed plots | 2 seedlings m-2 |
| Juvenile mortality in gaps | This study | Derived from statistical analyses |  |
| Juvenile mortality under closed canopy | This study | Derived from statistical analyses |  |
| Mature tree mortality | This study | Derived from statistical analyses |  |
| Annual mature tree growth rate <100years old | (Holzwarth *et al.* 2013) |  | 0.4 cm year-1 |
| Annual mature tree growth rate <100years old | (Holzwarth *et al.* 2013) |  | 0.35 cm year-1 |
| Annual mature tree growth rate <100years old | (Holzwarth *et al.* 2013) |  | 0.25 cm year-1 |
| Annual chance of mast year | (Packham *et al.* 2012) | In the UK beech trees mast once every three years on average | 0.3 |