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Museum specimens reveal loss of pollen host plants as key factor driving wild bee decline in The Netherlands

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Evidence for declining populations of both wild and managed bees has raised concern about a potential global pollination crisis. Strategies to mitigate bee loss generally aim to enhance floral resources. However, we do not really know whether loss of preferred floral resources is the key driver of bee decline because accurate assessment of host plant preferences is difficult, particularly for species that have become rare. Here we examine whether population trends of wild bees in The Netherlands can be explained by trends in host plants, and how this relates to other factors such as climate change. We determined host plant preference of bee species using pollen loads on specimens in entomological collections that were collected before the onset of their decline, and used atlas data to quantify population trends of bee species and their host plants. We show that decline of preferred host plant species was one of two main factors associated with bee decline. Bee body size, the other main factor, was negatively related to population trend, which, because larger bee species have larger pollen requirements than smaller species, may also point toward food limitation as a key factor driving wild bee loss. Diet breadth and other potential factors such as length of flight period or climate change sensitivity were not important in explaining twentieth century bee population trends. These results highlight the species-specific nature of wild bee decline and indicate that mitigation strategies will only be effective if they target the specific host plants of declining species.

Significance

Growing concern about bee declines and associated loss of pollination services has increased the urgency to identify the underlying causes. So far, the identification of the key drivers of decline of bee populations has largely been based on speculation. We assessed the relative importance of a range of proposed factors responsible for wild bee decline and show that loss of preferred host plant species is one of the main factors associated with the decline of bee populations in The Netherlands. Interestingly, species foraging on crop plant families have stable or increasing populations. These results indicate that mitigation strategies for loss of wild bees will only be effective if they target the specific host plants of declining bee species.


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sensitivity to climate change (4, 24, 25) may be associated with bee decline as well, and, to date, the relative importance of diet breadth and pollen host plant preference in explaining bee population trends remains unclear.

Here we solve this problem by analyzing historical pollen preferences of wild bees (15). Bees are generally more selective in their choice of food plants when foraging for pollen (source of protein and minerals for both larvae and adults) than nectar (source of energy) (26, 27). Distributional changes in plant species from which pollen is collected therefore probably exerts a larger influence on bee populations than changes in nectar plants. We investigate whether and to what extent loss of preferred floral resources drives bee population trends in The Netherlands, one of the most human-modified and intensively farmed countries in the world. Over the course of the twentieth century, agriculture has intensified in The Netherlands (Fig. S1) and the area of seminatural habitat preferred by bees has diminished to only one-fifth of the area at the beginning of the twentieth century (Fig. S2). More than half of the bee species are currently on the national Red List (28). As such, this country is a particularly suitable study area to identify critical factors associated with bee population decline.

We assessed pollen host plant use of bee species independently from their population trends by analyzing pollen loads on the bodies of bee specimens that were collected before 1950 (15), before the onset of agricultural intensification in The Netherlands. Altogether, our analysis included trend and trait data of 57 bee species in 10 genera and 4 subfamilies (Table S1). We calculated population trend indices for bee species and their host plants (period 1902–1949 vs. 1975–1999) using extensive national species distribution datasets (13, 29). Linear mixed models, with bee subfamily as a random factor to account for phylogeny, and a multi-model inference approach were used to examine the relationship between bee population trends and pollen host plant use, simultaneously taking into account differences in species’ rarity before the onset of agricultural intensification and other factors that have been proposed to explain bee population trends.

Results and Discussion

Model averaging across our set of candidate models (ΔAICc < 4) (30) revealed that population change of pollen host plants (full-model averaged standardized regression coefficient β = 0.54; relative variable importance ωp = 1.00), body size (β = −0.60; ωp = 1.00), and range size before 1950 (β = 0.20; ωp = 0.72) were the most important factors associated with bee population trends (Table 1). A model with just these three predictors best explained wild bee population change between the periods 1902–1949 and 1975–1999. It suggests that bee population trends were positively related with host plant change index (Fig. 1A) and initial range size (Fig. 1C) and negatively related with body size (Fig. 1B). This model explained 50% of the variation in bee population responses and had a probability of 0.37 of being the best model among the seven models in the candidate set. Analysis of bee trends based on a more extensive dataset that also included more recent bee records [period 1900–1989 vs. 1990–2011 (13)] produced similar results (Table S2). Our data do not enable us to distinguish whether the observed link between population trends of bees and their preferred host plants results from plant declines causing bee declines or vice versa. However, circumstantial evidence argues in favor of the proposition that it is primarily the loss of preferred host plants that is causing bee decline. First, distribution changes of plants in The Netherlands do not differ among insect-pollinated, wind- or water-pollinated, and self-pollinating plants (3), which reflects that loss of plant diversity in The Netherlands is mainly driven by abiotic factors associated with land use change, such as eutrophication, desiccation, and acidification (31, 32). Second, most insect-pollinated plants are pollinated by a diverse array of both generalist and specialist pollinators (33), which makes them fairly robust to the loss of a subset of pollinators. Bee losses may, for instance, have been compensated for by the increase of other important groups of pollinators such as hoverflies. Hoverflies, whose larvae do not depend on floral resources, have not been negatively affected by land use change and have even increased in The Netherlands over the last decades (3). Finally, a recent study on plant–pollinator networks in grasslands shows that land use intensity primarily drives loss of host plants, and that loss of host plants subsequently drives bee decline, whereas the reciprocal effects are not pronounced (34).

Length of flight period (β = 0.004; ωp = 0.15) and the degree of phenological advancement of the flight periods of bees between 1902–1949 and 1975–1999 (climate change sensitivity) (β = 0.001; ωp = 0.10) had little value in explaining bee population trends (Table 1). Mean daily temperatures during the activity period of bees (April–September) have increased in The Netherlands between 1906 and 2012 (Fig. S3). However, the extent to which bees advanced their flight periods in response to these rising temperatures did not contribute to explaining differences in bee population trends. This may indicate that bee species did not suffer from phenological mismatches with their host plants (35).

Surprisingly, diet breadth, i.e., the number of different pollen host plants used by a bee species, was also of minor importance in explaining bee population trends (β = 0.03; ωp = 0.25, Table 1). Species that use only a narrow array of food resources are generally expected to be more vulnerable to decline under environmental change (36). However, our results

### Table 1. Model selection and model averaging results for candidate models explaining bee population trends

<table>
<thead>
<tr>
<th>Predictor</th>
<th>Model 1</th>
<th>Model 2</th>
<th>Model 3</th>
<th>Model 4</th>
<th>Model 5</th>
<th>Model 6</th>
<th>Model 7</th>
<th>ΔAICc</th>
<th>ωm</th>
<th>95% CI</th>
<th>ωp</th>
</tr>
</thead>
<tbody>
<tr>
<td>Change of host plants</td>
<td>0.54</td>
<td>0.53</td>
<td>0.55</td>
<td>0.53</td>
<td>0.53</td>
<td>0.54</td>
<td>0.58</td>
<td>0.54</td>
<td>0.29</td>
<td>−0.79 to 1.00</td>
<td>1.00</td>
</tr>
<tr>
<td>Body size</td>
<td>−0.63</td>
<td>−0.62</td>
<td>−0.51</td>
<td>−0.51</td>
<td>−0.64</td>
<td>−0.63</td>
<td>−0.49</td>
<td>−0.60</td>
<td>−0.89</td>
<td>−0.31 to 1.00</td>
<td>1.00</td>
</tr>
<tr>
<td>Range size before 1950</td>
<td>0.29</td>
<td>0.25</td>
<td></td>
<td>0.30</td>
<td>0.28</td>
<td></td>
<td>0.40</td>
<td>0.20</td>
<td>−0.14</td>
<td>−0.54 to 0.72</td>
<td>0.72</td>
</tr>
<tr>
<td>Diet breadth</td>
<td>0.11</td>
<td>0.18</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>0.03</td>
<td>−0.14</td>
<td>−0.21 to 0.25</td>
<td>0.25</td>
</tr>
<tr>
<td>Length of flight period</td>
<td></td>
<td></td>
<td>−0.02</td>
<td>0.11</td>
<td></td>
<td>0.00</td>
<td></td>
<td>0.00</td>
<td>−0.11</td>
<td>−0.12 to 0.15</td>
<td>0.15</td>
</tr>
<tr>
<td>Phenological advancement</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>0.00</td>
<td></td>
<td></td>
<td>0.00</td>
<td>−0.07</td>
<td>−0.08 to 0.10</td>
<td>0.10</td>
</tr>
<tr>
<td>ωm</td>
<td>0.37</td>
<td>0.14</td>
<td>0.12</td>
<td>0.10</td>
<td>0.10</td>
<td>0.10</td>
<td>0.05</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Candidate models are ranked in order of increasing differences in corrected Akaike information criterion (ΔAICc). Explanatory variables were standardized by centering and dividing by 2 SDs. Akaike model weights (ωm) indicate the probability that a model is the best approximating model given the set of models considered. For each predictor, the parameter estimate for each candidate model is given, along with its model averaged estimate (β) (including zeros for variables that are not in a particular model), 95% confidence interval, and relative importance (ωp). Confidence intervals not overlapping zero are indicated in bold.
Relationship between bee population trends and body size, change index of pollen host plants, and initial rarity of bee species. Partial regression plots based on the best model in the candidate set for (A) log-transformed weighted mean change index of pollen host plants in bee species’ pollen diets, (B) body size measured as the intertuggular distance (ITD), and (C) log-transformed number of occupied \( 5 \times 5 \) km grid cells before 1950. Note that a bee change index of 1 indicates no change. Plotted points represent partial residuals. Shaded areas indicate 95% confidence bands.

The observed negative relationship between bee body size and population trend may result from the decline of the large-bodied bumblebee species, which as a group have experienced particular strong declines in Europe (13, 40). Yet, body size remained a key factor (\( \beta = -0.39; P = 0.93 \) even when the 10 bumblebee species were excluded from the analysis, suggesting that the negative relationship with body size does not only reflect the decline of the bumblebees. Alternatively, the effect of body size may be linked to loss of floral resources as driving factor for wild bee decline. Large bee species may be more susceptible to land use change than smaller ones (4, 23) because of their larger pollen quantity requirements (41). In homogenous, intensively farmed landscapes, declining floral resources may cause fewer problems for small species to find sufficient food for offspring production than for large species, despite the larger foraging ranges of large species (42).

To identify whether bee decline was associated with preference for certain plant families, we calculated for each plant family the mean trend index of bees that relied for at least 10% of their pollen diet on that particular plant family (Fig. 3). On average, bee species that preferentially collected pollen from plant species belonging to the Rosaceae family significantly increased while bees feeding on plants in the Fabaceae family decreased. This pattern remained when threshold values of 20% or 30% were used (Fig. S4). The declines in Fabaceae species can be attributed to the loss of Fabaceae-rich seminatural grasslands (14), but also to the drastic decrease of the agricultural use of Fabaceae as fodder and cover crops during the twentieth century (Fig. S5). Our findings shed new light on the consequences of reported losses of wild bees for crop pollination. Bees that preferentially collect pollen from Rosaceae, Brassicaceae, and Asteraceae species, which contain major insect-pollinated crops such as apple, strawberry, oil-seed rape, and sunflower, have remained stable or actually increased. Delivery of crop pollination services by wild bees may therefore be under less threat from land use change pressures than preservation of wild bee diversity.

Because it is impossible to experimentally examine causes of large-scale population declines, it is difficult to establish causal relationships between drivers and population trends. The robust linkage of bee population trends, through historical host plant preferences, to plant population trends that was found in this study therefore probably represents the best possible evidence for key factors causing bee population decline in intensively farmed landscapes for some time to come. These insights will help us develop more effective ways to mitigate loss of species that are threatened in their existence as well as promote bees important for delivery of crop pollination services.

Methods

Examined Bee Species. In The Netherlands, 357 wild bee species are found. A total of 256 species actively collect and transport pollen to provision their offspring. The remaining bees are cleptoparasitic bees that lay their eggs in brood cells of host bees and do not forage for pollen themselves. We focused our sampling efforts on bee species that use multiple host plant species (“polylectic bee species”), which constitute 70% of the pollen-collecting bee species in The Netherlands (13). We used the number of \( 5 \times 5 \) km grid cells occupied before 1950, obtained from the national bee distribution database.
(13), to assess species’ rarity before the onset of major environmental change, and only included species if they were common (present in at least one-hundred-fifty 5 × 5 km grid cells), fairly common (70–149 grid cells), or only moderately rare (20–69 grid cells) before 1950. Bee species from the genus Hylaeus were excluded because female bees of these species transport pollen internally in their crops rather than externally on their bodies, which makes nondestructive sampling of pollen difficult. This resulted in a total of 75 bee species for which we aimed to determine pollen host plant use (see Pollen Host Plant Use).

Bee Population Trends. Relative bee population trends were determined using the national bee distribution database of European Invertebrate Survey (EIS) - The Netherlands (13). This database contains 186,147 records of bees collected and observed in The Netherlands between 1809 and 2011. Relative change indices for the focal bee species were calculated as the ratio between the number of occupied 5 × 5 km grid cells in the period 1902–1949 (26,749 records) and the number of occupied grid cells in the period 1975–1999 (45,447 records), divided by the average of the change ratio of all focal bee species. Relative change indices thus reflect bee species’ performance between the two periods relative to the average performance of all of the species considered. We only used data of grid cells that were inventoried in both periods (total of 568 cells) and, to avoid potential bias arising from the much larger proportions of field observations in the database in recent years (which are mainly restricted to common and easily recognizable species), only included records from natural history collections. Sampling intensity differed between periods, which, as the probability that a species is recorded depends on sampling intensity, may introduce bias when comparing grid cell trends between both periods. To compare both periods, we used the total number of bee records per period as a proxy for sampling intensity, we corrected for differences in sampling intensity between periods by multiplying the number of grid cells each species occupied in 1975–1999 with the quotient of the total number of bee records in 1902–1949 divided by the total number of bee records in the 1975–1999 period (13). The rationale behind comparing the 1902–1949 and 1975–1999 periods is that these periods form the basis for the national Red List of terrestrial plants (29) (Bee Population Trends) and contrast a period of low-intensity farming with a period of intensive agriculture and reduced availability of seminatural bee habitat (Figs. S1 and S2). However, to assess whether the results of our study were robust to the chosen time periods, we also analyzed the drivers of wild bee decline using bee trends based on the periods used by Reemer et al. (13) [1900–1989 (77,920 records) vs. 1990–2011 (68,491 records), total of 858 grid cells]. Using trends based on these time periods produced results similar (Table S2) to the results reported in the main text (Table 1).

Pollen Host Plant Use. Pollen loads of female bees of the selected species were sampled in the entomological collections of the Natural History Museums of Amsterdam, Leiden, Leeuwarden, Rotterdam, Tilburg, Wageningen, and Brussels (Belgium). Samples were only taken from specimens that were collected before 1950 (between 1870 and 1950) and that had pollen in the pollen-carrying bodily structures (scopa or corbicula). Samples of pollen from each pollen load were mounted in glycerine jelly containing basic fuchsin to stain the pollen grains. Identification was done using a light microscope at 400× magnification with the assistance of a reference pollen collection of ~130 species and reference documents (43). Pollen grains were identified to the lowest possible taxonomic level, mostly to genus (67%). For each sample, we estimated the percentage contribution of each pollen taxon, with pollen taxa contributing <5% not being considered as they may result from contamination. To avoid potential bias resulting from several samples collected at the same location at the same date, duplicate samples were randomly excluded from the dataset.

We could not reliably determine pollen host plant use for 18 of the selected 75 species because of insufficient numbers of pollen samples (n < 15) in museum collections, and these species were omitted from further analyses. The mean number of pollen samples per species of the final set of 57 bee species was 28.9 (min = 15, max = 63). These 57 species (Table S1) represent 40% of the pollen-collecting bee species that were observed in at least twenty 5 × 5 km grid cells in The Netherlands before 1950. The majority of the specimens of these species were collected at locations in The Netherlands, but a limited number (7%) were collected at Belgian locations near the Dutch–Belgian border. Out of the total number of 1,646 specimens, broadly equal numbers of specimens were sampled as part of the study area (respectively 863 and 783). In total, we identified 170 different pollen taxa in the pollen loads of the 57 focal bee species.

Following Klein and Raemakers (15), we quantified pollen host plant preference for each bee species as the percentage contribution of each...
pollen taxon to the total pollen load sampled from all specimens of the bee species. Population changes of pollen host plants during the twentieth century were based on the frequency of occurrence of plant species in 1 x 1 km grid cells in the periods 1902–1949 (1.7 million records) and 1975–1999 (3.5 million records) (29, 44). We calculated relative change indices of host plants as the ratio between the number of occupied grid cells in the period 1902–1949 and the number of occupied grid cells in the period 1975–1999, divided by the average of the change ratio of all identified bee host plants. Change indices were based on a selection of 7,374 grid cells with multiple observations within the grid cell across both periods (nearly 25% of the land surface of The Netherlands), corrected for temporal differences in sampling intensity (29). For pollen taxa that could only be identified to genus or family level, we used distribution data of the common plant species in the respective genus or family to calculate a weighted average change index for these pollen taxa (15).

Next, for each bee species, we calculated the weighted mean population change index of the host plants in its pollen diet \( C_{\text{pollen}} \) as

\[
C_{\text{pollen}} = \frac{\sum_{i=1}^{n} PC_i \times CI_i \times \omega_{p,m}}{\sum_{i=1}^{n} PC_i}
\]

with \( PC_i \) representing the percentage contribution of pollen taxon \( i \) to the total observed pollen load of the bee species, \( CI_i \) the relative change index of pollen taxon \( i \), and \( n \) the total number of pollen taxa observed in the pollen loads of the bee species. Pollen diet breadth, i.e., the number of pollen taxa in the pollen diets of bee species, was determined after sample-based rarefaction to 15 samples using EstimateS software (45).

Other Factors Associated With Bee Decline. For each bee species, we assessed its initial rarity, body size, phenology, and response to climate change. First, as rarity in itself may be an important cause of population decline (18) and may confound any observed relationship between bee traits and population trends (17), we quantified species’ rarity before the onset of major land use changes in The Netherlands as the number of 5 x 5 km grid cells before 1950.

Second, we measured bee body size as the interangular distance (ITD). The ITD is the distance between the two insertion points of the wings, which is a reliable estimator of bee body size (46). For all species, except Bombus species, we measured the ITD of 10 female specimens. For Bombus species, we measured the ITD of 20 worker bees to account for larger intraspecific variation in body size in these species.

Third, we determined the start and length of the flight periods of bees using the national bee distribution database of EIS - The Netherlands (13). For both the 1902–1949 and 1975–1999 periods, we used the records in the bee database (mean number of records per species 1902–1949 = 92; 1975–1999 = 374) to calculate the 10th and 90th percentile of the recording day (1 January = 1) for each bee species and defined the 10th percentile as the start, and the number of days between the 10th and 90th percentiles as the length of the flight period of bee species (4). Seven species had become too rare (less than 40 records) to reliably determine their flight periods in the period 1975–1999. For these species, we predicted the start and length of the flight period using the linear relationships between the other species’ start of flight period \( \text{Start}_{1975-1999} \) = 0.696 \times \text{Start}_{1902-1949} + 14.629, \( F_{1,49} = 484.1, P < 0.001, R^2 = 0.91 \) and end of flight period in 1902–1949 and 1975–1999 \( \text{End}_{1975-1999} = 0.920 \times \text{End}_{1902-1949} + 15.929, F_{1,49} = 421.7, P < 0.001, R^2 = 0.90 \). We quantified the influence of climate change by calculating the phenological advancement of bee species’ flight periods between both periods \( \text{Start}_{1975-1999} - \text{Start}_{1902-1949} \).

Although nesting ecology may be expected to be an important factor underlying different responses of bee species to land use change (11, 47), still little is known about the nesting requirements of most bee species, and nesting ecology is difficult to quantify. Rough categorical classifications of bee species’ nesting ecology (e.g., nesting above ground vs. nesting below ground) oversimplify the broad array of nesting habitats, substrates, and construction materials used by different bee species (48) and probably obscures any relationship between bee species’ nesting ecology and population response to land use change. Possibly as a consequence, previous analyses did not find any relationship between nesting ecology and population trends of species (46). We therefore chose not to include any categorical measure of nesting ecology in our analysis.

Floral Resource Availability in Contemporary Agricultural Landscapes. In 2012, we used a stratified sampling approach to estimate spring and summer floral resource availability in 16 agricultural landscapes (1 km radius) in The Netherlands. Flower inventories were conducted in habitats in seven main land use classes: seminatural habitats (e.g., forest edges, wooded banks), cultivated grasslands, nonflowering crop fields (e.g., maize fields, wheat fields), flowering crop fields (oilseed rape fields), uncultivated field boundaries (field boundaries, ditch banks, road verges), nonflowering perennial habitats (forest interiors and gardens). Flower inventories were performed twice: once in May and once in July. For each of the land use classes, we estimated percentage cover of flowering forbs in randomly selected plots (100 m²), evenly spread out over the study landscapes (grand total of 48 plots). We used GIS to calculate the area of land use classes in the 1-km radius study landscapes, and calculated the estimated total landscape-wide floral resource availability (% cover) in May and July in each landscape as: (mean flower cover land use class A \( \propto \) proportion of land use class A in the study landscape + mean flower cover land use class B \( \propto \) proportion of land use class B in the study landscape + …) / 100.

Data Analysis. We used linear mixed models and an information theoretic approach to assess to what extent the explanatory variables were related to bee population trends. We constructed a set of linear mixed models containing all possible combinations of the different predictors, including an intercept only model. As phylogenetic analyses are controversial (49) and phylogenetic trees for bees are continuously challenged (50), we did not use phylogenetic regression but instead included bee subfamily as a random factor to account for potential nonindependence of trends and traits among closely related bee species (4). The variables “range size before 1950” and “change of host plants” were log-transformed to reduce positive skew. We excluded start of flight period (correlated with change of host plants, \( r = -0.53, P < 0.001 \)) from the set of predictors to avoid multicollinearity. A multimodel inference approach that used start of the flight period instead of change of host plants showed that the highest-ranking model that included start of flight period (AICc = 90.3, Akaike model weight \( \omega_{p,m} = 0.016 \)) was 62 times less likely to be the best model than the highest-ranking model that included change in host plants (AICc = 82.0, \( \omega_{p,m} = 0.984 \)).

We ranked the 64 possible models according to their Akaike information criterion corrected for small sample size (AICc) (Table S3) and restricted our candidate model set to models with a AICc < 4 (30). For each model in the candidate set, we calculated the Akaike model weight \( \omega_{p,m} \), which reflects the probability that a model is the best approximating model given the set of candidate models considered (30). Marginal \( R^2 \) values (the variance explained by the fixed effects variables) of models were calculated following Nakagawa and Schielzeth (51). To account for model selection uncertainty, we calculated full-model averaged parameter estimates (including zero when predictors were not included in a particular model (52)) for each predictor in the candidate model set. The relative importance \( \omega_{0.05} \) of a predictor was based on the sum of the Akaike weights across all models in the candidate model set that included the predictor (30). All analyses were performed using R (53), using the packages lm4 (54) and MuMln (55).

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