The effect of spatial variation for predicting aphid epidemics

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3500 words, 2 tables, and 3 figures.

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Abstract

In order to improve forecasting of aphid epidemics, it is important to know the spatial scale at which specific forecasts are reliable. To investigate the spatial scale of aphid epidemics, we have developed a spatio-temporal stochastic aphid population growth model, and fitted the model to empirical spatial time-series aphid population data using a Bayesian hierarchical fitting procedure. Furthermore, detailed spatial data of the initial phases of epidemic development was investigated in a semivariogram. Our results suggest that there is limited spatial variation in the initial occurrence probability at a spatial scale of 10 km. Consequently, the results support the hypothesis that initial aphid population sizes and epidemics may be predicted in fields within a 10 km radius. For farmers, this may imply that they can rely their decision of whether to spray against aphids on observations made by other nearby farmers or by the consultancy service.

Keywords: Aphid population model; spatial and temporal model; hierarchical models

Introduction

The common cereal aphids, the Cherry oat aphid (*Rhopalosiphum padi*), the Grain aphid (*Sitobion avenae*) and the Rose-grain aphid (*Metopolophium dirrhodum*), are known to cause considerable losses to winter wheat in large parts of the world, and these species have been estimated to cause losses of 700.000 tons per year in Europe due to direct damage (Wellings et al. 1989). On top of this figure comes the indirect damage due to virus transmission and sooty mold caused by the excretion of honeydew (Larsson 2005). Therefore, there is a great interest in controlling aphids in winter wheat, and a number of simulation models have been developed to predict the population development of these pests (e.g. Ciss et al. 2014; Duffy et al. 2017; Holst & Ruggle 1997; Klueken et al. 2009). Furthermore, some population dynamic models have been combined with modules on economy and pesticide choices to produce decision support systems with the aim of helping farmers optimizing the timing of pesticide applications. Such decision

support systems include CPO in Denmark (Hagelskjær & Jørgensen 2003) and GETLAUS (Gosselke et al. 2001) in Germany. In a review of decision support systems, (Axelsen et al. 2012) concluded that farmers generally do not use decision support systems, and one of the reasons is that they do not find it worth the effort to spend time on estimating aphid input densities. Instead, many farmers perform precautionary insecticide treatments, which may not be economically sound.

In order to optimize the decision support systems, large efforts have been spent on improving model performance in relation to weather parameters and natural regulation etc. These efforts have without doubt improved the models, but taking into consideration that aphid populations develop exponentially, and that the population can double in 55 hours at 20°C (Dedryver et al. 2010), it is critical that the initial densities are determined at high precision in order to predict the epidemic development. However, surprisingly little efforts have been devoted to obtaining quick and reliable estimates of the initial densities, although several methods have been developed to estimate the density of aphids in cereals. For instance, Elliott et al. (1990) developed binomial sequential sampling plans that require rather larger efforts at low densities to produce reliable estimates, and Hansen (1991), who suggested to investigate aphid presence/absence on 50 or 100 tillers and used an equation to convert to aphids per tiller. Both methods take some time, and come up with assessments of average densities with some uncertainties. The uncertainties are predefined in the sequential sampling plan and the required uncertainty level is decisive for the number of plants to investigate. When counting presence/absence on a number of straws, the uncertainty can be calculated before being used in simulation models and decision support systems. However, none of the models and decision support systems appear to use uncertainty in their projections of aphid population development, and in turn relate uncertainties to the output. Given that aphids show exponential population growth, uncertainties in the estimate of initial densities can cause large uncertainties to the projections of the aphid population density some weeks later. This uncertainty should ideally be reflected in the suggestions produced by decision support system.

To predict the spatial and temporal development of aphid epidemics it is important to sample and model population data that encompasses both spatial and temporal dimensions. In this study, we have sampled aphid populations in a spatial setup during both the initial and the epidemic phase to fit an epidemic model. More specifically, we have developed a spatio-temporal stochastic aphid population growth model and fitted the model to empirical spatial time-series aphid population data using a Bayesian hierarchical fitting procedure. Such Bayesian hierarchical population models have successfully been applied on a number of pest cases, e.g. the population dynamics of coffee berry borer infestation (Ruiz-Cárdenas et al. 2009). The fitted spatio-temporal population growth model may be used to generalize existing deterministic aphid forecasting models with the effect of stochastic spatial variation. Furthermore, we use the fitted model and complementary spatial statistics to investigate the hypothesis that initial aphid population sizes and epidemics may be predicted in fields within a 10 km radius of the nearest aphid-monitoring site.

Materials and Methods

Field sites and aphid sampling

The occurrence probability of the grain aphid, *Sitobion avenae*, was recorded at twelve wheat fields in the middle of Jutland, Denmark, in 2016 and 2017. The wheat fields (=sites) were laid out in hierarchical geographic design with three regions of four sites. Each region had a center site with sites positioned approximately three, six and ten kilometers away (Fig. 1). At each site the occurrence probability of aphids on individual plants were recorded and the occurrence probability was used as a proxy for the aphid population size (Hansen 2003).

.In 2016, aphid occurrence was recorded for five samples of either 80 or 100 wheat plants at each site on May 24, June 6, June 13, June 20, June 27, July 4, and July 11. The five samples within a site were taken along an irregular transect with at least 50m between plots. Site-specific degree-days were calculated from

the average day temperature at the weather station closest to each site with a base temperature of 5 degrees Celsius. Furthermore, the intensity of precipitation events at the different sites was recorded.

In order to complement the result of the spatial variation of the initial phases of aphid epidemics obtained in the spatial modelling of the aphid occurrence data sampled in 2016, aphid occurrence was recorded more intensively at the same twelve sites in the beginning of the growth period the following year and analyzed in a semivariogram. In 2017, aphid occurrence was recorded for ten samples of 50 wheat plants at each site on May 30 and June 7. The ten samples within a site were laid out along three transect with at least 50m among all plots, and the exact geographical position of each sample was determined.

Statistical modelling of the aphid population

The spatio-temporal aphid occurrence data is modelled using Bayesian hierarchical methods (Clark & Gelfand 2006). The observed number of straws with at least one aphid at site *i* and plot *k* at degree-day *t* is denoted $y_{i,k,t}$ and is assumed to be binomially distributed with $n_{i,k,t}$, the number of straws sampled, and $p_{i,t}$, the occurrence probability that a straw has at least one aphid at site *i* at degree-day *t*,

$$y_{i,k,t} \sim Bin(n_{i,k,t}, p_{i,t}) \quad (1).$$

The site-specific occurrence probability is modelled using an exponential function of degree-day *t*,

$$p_{i,t} = p_{i,0} Exp(r_0 + r_1\, t + r_2\, t^2 + \epsilon_i) \quad (2),$$

where $p_{i,0}$ is the occurrence probability on a fixed initial day, r_0, r_1, and r_2 are population growth parameters, and ϵ_i are Gaussian distributed site-specific random effects, $\epsilon_i \sim Gau\ (0, \sigma_p^2)$.

The *n* site-specific initial occurrence probabilities are assumed to arise from a Gaussian process model,

$$p_{i,0} \sim Gau\ (\mu_0, \Sigma),\ \Sigma = \begin{pmatrix} \sigma_0^2 & \cdots & \sigma_0^2\ \rho(d_{1,n}) \\ \vdots & \ddots & \vdots \\ \sigma_0^2\ \rho(d_{n,1}) & \cdots & \sigma_0^2 \end{pmatrix} \quad (3),$$

where μ_0 is the mean initial occurrence probability, σ_0^2 is the variance, and $\rho(d_{i,j}) = \rho_0\ Exp(-\frac{d_{i,j}}{\alpha})$ with $d_{i,j}$ being the distance between site *i* and site *j* , α is the scale of the spatial effect that is set to 10 km, and ρ_0 is a parameter that measures the spatial covariance (Haran 2011; Ovaskainen et al. 2016). The covariance matrix by definition has to be positive definite, which puts upper and lower bounds on ρ_0.

The prior distributions of all parameters were assumed to be uniform except the two scale parameters that were assumed to be inverse gamma distributed, $\sigma_p \sim Invgam(0.1,0.1)$ and $\sigma_0 \sim Invgam(0.1,0.1)$. The joint Bayesian posterior distribution of the parameters and the two latent variable vectors, $p_{i,0}$ and ϵ_i, each of dimension *n*, were calculated using Markov Chain Monte Carlo (MCMC), Metropolis-Hastings, simulations with a multivariate normal candidate distribution and using a MCMC run of 300,000 iterations after a burn-in period.

Plots of the sampling chains of all parameters and latent variables were inspected in order to check the mixing properties of the used sampling procedure, and whether the burn-in period was sufficient. Additionally, the overall fitting properties of the model were checked by inspecting the regularity and shape of the marginal distribution of all parameters as well as the distribution of the deviance ($= -2 \log L(Y|\theta)$). The efficiency of the MCMC procedure was assessed by inspecting the evolution in the deviance.

Statistical inferences on the parameters were based on the marginal posterior distribution of the parameters.

All calculations were done using *Mathematica* version 10 (Wolfram 2015).

Results

The observed spatio-temporal mean occurrence probability of *S. avenae* in 2016 (Fig. 2) was fitted to the spatial growth model. The burn-in period of the MCMC was relatively long (600,000 iterations), but after the deviance had stabilized, the fitting properties of the models were judged to be acceptable based on visual inspections of the mixing properties of the parameters and the latent variables.

The marginal posterior distributions of the parameters of interest are summarized in Table 1 by their percentiles. The parameters were generally uncorrelated, except for, as expected, the estimates of the population growth parameters r_0, r_1 and r_2, which were highly correlated (Table 2).

The deterministic part of the population growth model (eqn 1.) seemed to adequately model the dynamics of aphid occurrence probabilities as a function of degree days from June 1 2016 to July 1 2016 (Fig. 3) when shape and mode of the expected epidemic was visually compared to the observed spatio-temporal mean occurrence probability data in the same period (Fig. 2).

The posterior marginal distribution of the parameter that measures the effect of geographic distance on the spatial covariance, ρ_0, is left-skewed towards the upper boundary and significantly larger than zero (Table 1), and the site-specific initial occurrence probabilities are consequently positively correlated among the sites at the spatial scale of 10 km. However, the importance of this positive correlation for the among-site variation in aphid epidemics has to be evaluated in relation to the estimated among-site variation in the initial occurrence probability as modelled by σ_0 , and population growth as modelled by σ_p (Table 1).

It has been hypothesized that episodes of heavy rain may lead to decimation of the aphid population. However, if the observed rain records at the sites, which included several episodes of heavy rain, were manually scored according to severity and compared to the latent variables that model the among-site

variation in population growth, ϵ_i, then there was no significant relationship between the heavy rain score and a relatively low population growth rate (P = 0.89).

The variation in aphid occurrence among sampling plots as a function of the geographical distance among plots on May 30 and June 7 2017 is shown as a semi-variogram (Fig. 4). Generally, the variation among plots is relative low at the two sampling days, although the variation seems to increase irregularly with time when assessed from only two samples in time. There is a slight indication that the variation among plots increases with the distance among plots at the second sampling.

Discussion

The site-specific initial occurrence probabilities was found to be positively correlated among the sites at the spatial scale of 10 km. However, when the spatial variation in the initial occurrence probability was examined in more detail the following year, the spatial variation among plots in the beginning of the aphid epidemics did not seem to increase much with among-plot distance. This indicates that there was only limited spatial effects, i.e. that the initial epidemic development was more or less in synchrony over distances up to and above 10 km. Since the parameter that measures the effect of geographic distance on the spatial covariance, ρ_0, also depends on the spatial variation in the following aphid epidemic estimated from the 2016 data set, we tend to put more weight on the more detailed investigation in 2017, and conclude that our investigation suggest that there is limited spatial variation in the initial occurrence probability. Consequently, the overall results support the working hypothesis that initial aphid population sizes and epidemics may be predicted in fields within a 10 km radius of the nearest aphid-monitoring site. For farmers, this may imply that they can rely their decision of whether to spray against aphids on observations made by other nearby farmers or by the consultancy service.

We did not detect any significant effects of heavy rain events on aphid occurrence probability. This is contradictory to the findings by Mann et al. (1995) that heavy rainfall may cause a loss of aphids up to 30 % of the population, but the recorded rainfall events in our fields may have been relatively benign.

The simple quadratic function used to model population growth as a function of degree-days performed adequately for modelling the growth in the aphid population from June 1 2016 to July 1 2016. However, the purpose of this simple quadratic model was only to model the deterministic part of the population growth in order to quantify the stochastic variation among sites, and the fitted quadratic model is not suitable for making actual predictions outside the domain of the collected aphid population data. Instead, the deterministic part of the population growth could be the models described by Holst et al. (1997), Ciss et al. (2014), Duffy et al. (Duffy et al. 2017) or the model element of the aphid modules of decision support systems such as CPO (Hagelskjær & Jørgensen 2003) and GETLAUS (Gosselke et al. 2001).

Most decision support systems will let the user know if it is appropriate to treat against aphids or not, without telling how certain the “decision” is. If the output came with uncertainties, such as “It can now with 55% certainty pay off to treat against aphids in your field”, using the decision support system may appear more difficult, because the farmer is left with an uncertain foundation for his decision. However, the output might include information on how to reduce the uncertainty. If the farmer had counted aphids on 50 tillers, the output might tell the farmer that the uncertainty would be reduced if he continued and counted on a higher number of tillers, and he could continue until he was able to take certain decisions. This would require some time, but the farmers would get some knowledge on the importance of spending time on providing the decision support system with good initial aphid density estimates, as spending more time will provide a safer foundation for decisions. Estimates of initial densities can be used to predict peak densities of *S. avenae* (Honek et al. 2017), and good estimates of initial densities will, everything else equal, provide better estimates of peak densities.

Nevertheless, the farmer might still not be willing to spend the required time on counting aphid, but counting might be found worth the effort if the result could be used by neighbors, or by all farmers within a certain area. This would optimize the balance between time input spend on counting aphids and economic output in terms of higher yield and probably less expenses on pesticide applications.

Tables

Table 1. Marginal posterior distributions of the parameters in the spatio-temporal stochastic aphid population growth model spatial when fitted to 2016 occurrence data. The distributions are summarized by their 2.5%, 50% and 97.5% percentiles.

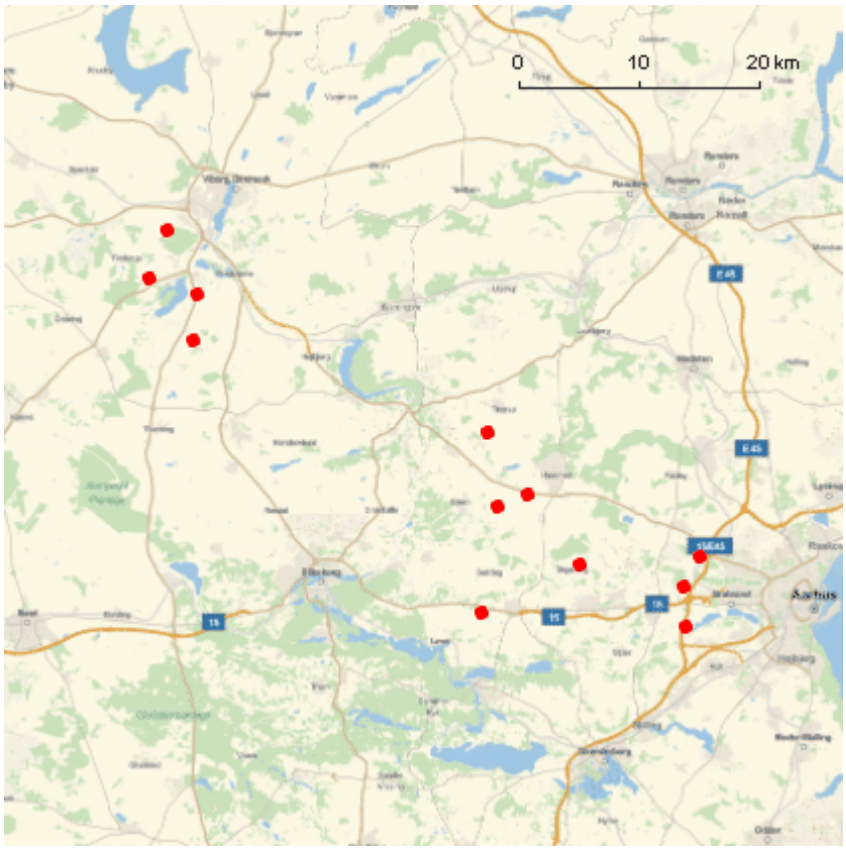
Parameter	2.5%	50%	97.5%
r_0	0.0147	0.0175	0.0215
r_1	-1.17E-05	1.32E-05	2.87E-05
r_2	-1.17E-07	-8.83E-08	-4.71E-08
σ_p	0.00554	0.00911	0.01677
μ_0	0.000395	0.00786	0.01531
σ_0	0.00509	0.00857	0.01620
ρ_0	0.936	1.2185	1.2412

Table 2. Correlation matrix between the parameters r_0, r_1, and r_2 in the spatio-temporal stochastic aphid population growth model spatial when fitted to 2016 occurrence data.

	r_0	r_1	r_2
r_0	1	-0.96	0.93
r_1	-0.96	1	0.99
r_2	0.93	0.99	1

Figures

Fig. 1. Twelve wheat fields in the middle of Jutland, Denmark, where aphids were sampled. The wheat field were laid out in hierarchical geographic design with three regions of four fields. Each region had a center field with fields positioned approximately three, six and ten kilometers away.



207 Fig. 2. Observed spatio-temporal mean occurrence probability of *S. avenae.* The sites are located in three

208 different regions (Borum, Hammel and Viborg). Each region had a center field (0) with fields positioned

209 approximately three (3), six (6) and ten (10) kilometers away.

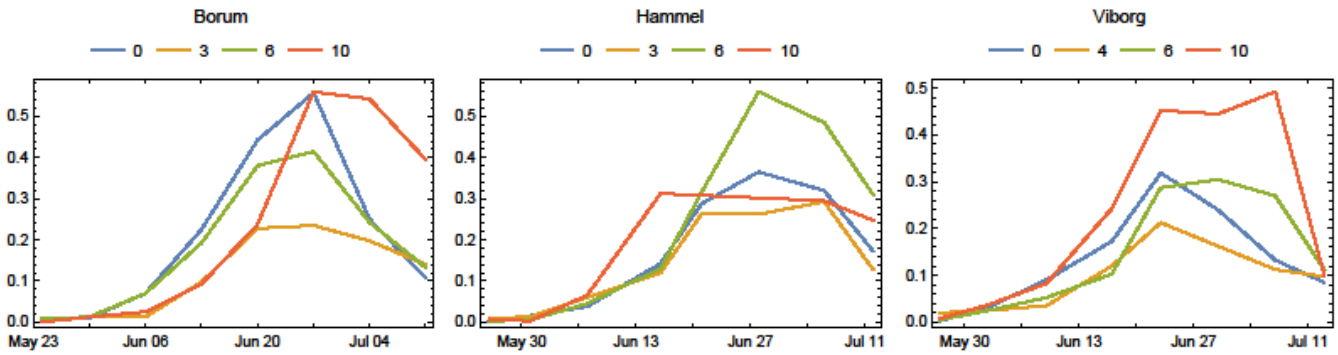


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Fig3. Expected occurrence probability of *S. avenae* at the mean degree-days from June 1 2016 to July 1 2016. The curve is calculated using equation (2), the expected μ_0, and the joint posterior distribution of r_0, r_1, and r_2. The thick line is the median occurrence probability and the two thin lines are the 25% percentile and 75% percentile of the occurrence probability. The shaded area is the 95% credibility interval of the occurrence probability.

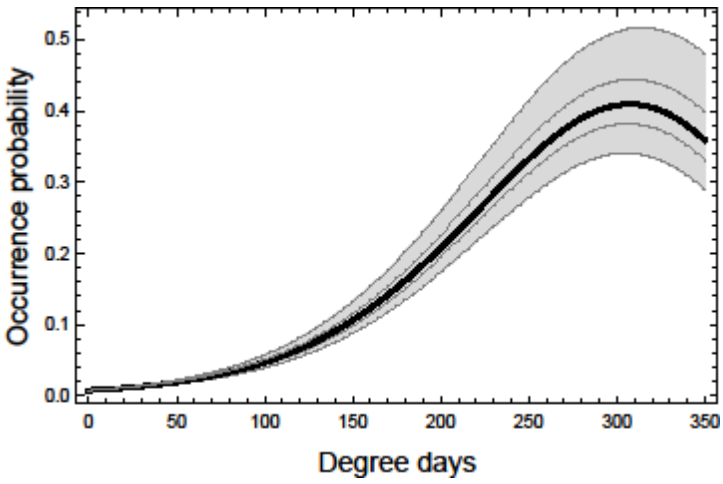
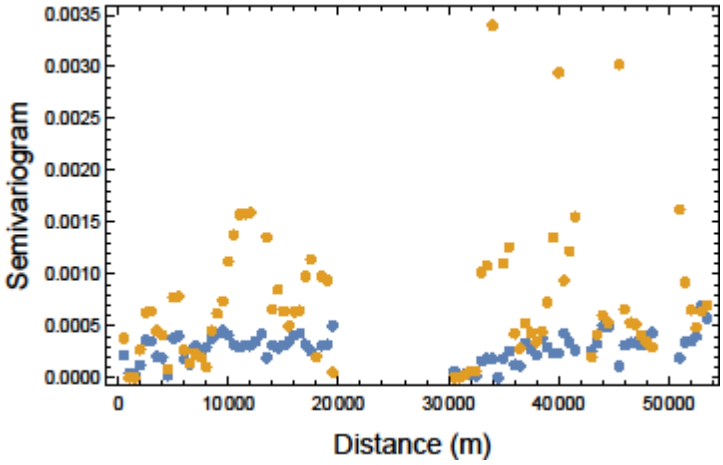


Fig. 4. Semivariogram. The variation in aphid occurrence as a function of the geographical distance among sampling plots. Blue points: May 30 2017, yellow points June 7 2017.



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