Impact of fungi on control of bollworms Chrysodeixis includens and Helicoverpa armigera

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22nd November 2018

VIII Encontro dos Alunos em Estatística e Experimentação Escola Superior de Agricultura Luiz de Queiroz

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### Outline

- 1. Introduction
- 2. Models and methods
- 3. Results and discussion
- 4. Final remarks

# <sup>1</sup> Introduction

### **Entomological motivation**

#### **Species of interest:**

- Chrysodeixis includens: the "soybean looper";
- Helicoverpa armigera: the "cotton bollworm";
- They feed feed on a wide range of plants, including many important cultivated crops (agronomic crops: soybean, cotton, maize, etc. and also vegetable and floricultural crops).

#### **Biological pest control:**

- Controlling pest population using other organisms;
- An important biological control agents are pathogenic fungi.

#### **Research question:**

The inoculation of fungi in soybean plants may inhibit the development of bollworms *Chrysodeixis includens* and *Helicoverpa armigera*?

# Design of the experiment

#### Treatments:

Three species of fungi:

*Metarhizium anisopliae* ESALQ-1638 (Met 1638); *Beauveria bassiana* ESALQ-3399 (Bb 3399); and *Isaria fumosorosea* ESALQ-3422 (If 3422);

Control (Tween 80).

#### **Experiment with whole plants**:

- The fungi were inoculated on the commercial subtracts to cultivate the soybean plants.
- 30 bollworms (for each treatment) were confined in a pot with a plant, where the substrate was isolated.
- The plots were evaluated every three days during 18 days for *Chrysodeixis includens* and 21 days for *Helicoverpa armigera*.
- This design were repeated two times in different periods.

**Outcomes:** 

- Weight of bollworms over time (longitudinal data) and
- Time to death of the bollworms (time-to-event data).

# Design of the experiment



Figure: Pictures of the experiment: (a) *Helicoverpa armigera*, (b) soybean plants, (c)-(b) bollworms confined in the pots.

### **Descriptive analysis (longitudinal data)**



Figure: Data on weights of the *Chrysodeixis includens* bollworms over time. The symbols ● and ▲ indicate death and pupa stage, respectively.

### Descriptive analysis (longitudinal data)



Figure: Data on weights of the *Helicoverpa armigera* bollworms over time. The symbols ● and ▲ indicate death and pupa stage, respectively.

### **Descriptive analysis (time-to-event data)**



Figure: Kaplan-meier survivor function estimates for the times to death of the bollworms.

# <sup>2</sup> Models and methods

## Non-linear models (logistic growth)

To model the weight growth of bollworms (*Y*) over time (*t*), we consider the logistic growth,

$$E(Y) = f(t) = \frac{\theta_{\rm A}}{1 + \exp[(\theta_{\rm M} - t)/\theta_{\rm S}]},$$

where

- ▶  $\theta_A$  is the horizontal asymptote (f(t) when  $t \to \infty$ , if  $\theta_S > 0$ ),
- $\theta_{M}$  is the inflection point of the curve and
- $\theta_{\rm S}$  is the scale parameter.



### Heteroscedastic non-linear mixed models

#### Statistical challenges

- Model the correlation between measures of the same bollworm and
- Model the heteroscedastic within-error.

### Fitted model

Let (y<sub>ijk</sub>, t<sub>ijk</sub>) denote the weight and time (days) of the *i*-th bollworm on the *j*-th treatment at the *k*-th time, the fitted model can be expressed as

$$y_{ijk} = \frac{\theta_{\mathrm{A}j} + b_{\mathrm{A}i}}{1 + \exp[(\theta_{\mathrm{M}j} + b_{\mathrm{M}i} - t_{ijk})/\theta_{\mathrm{S}j}]} + \varepsilon_{ijk},$$

#### Variance components

$$\begin{bmatrix} b_{\mathrm{A}i} \\ b_{\mathrm{M}i} \end{bmatrix} \sim \mathcal{N}\left( \begin{bmatrix} 0 \\ 0 \end{bmatrix}, \mathbf{\Sigma} = \begin{bmatrix} \sigma_{\mathrm{A}}^2 & \sigma_{\mathrm{AM}} \\ \sigma_{\mathrm{AM}} & \sigma_{\mathrm{M}}^2 \end{bmatrix} \right),$$
$$\forall \operatorname{Var}(\varepsilon_{ijk}) = \sigma^2 \delta_{1j}^2 \delta_{2K}^2.$$

### Accelerated failure time models

Let  $T_{ij}$  be the time-to-death of the *i*-th treatment and *j*-th bollworm, the AFT model can be expressed (where  $\omega = \log(\varepsilon)$ ) as

$$T_{ij} = \exp(\boldsymbol{x}_{ij}^{\top}\boldsymbol{\beta})\boldsymbol{\varepsilon}^{\sigma}$$
$$\log(T_{ij}) = \boldsymbol{x}_{ij}^{\top}\boldsymbol{\beta} + \sigma\boldsymbol{\omega}.$$

The distributional assumption of  $T_{ij}$  implies distribution of  $\omega$  (e.g.  $T_{ij} \sim \text{Weibull}(\alpha, \lambda) \implies \omega \sim \text{E.V.}(\lambda, \alpha)$ ).

#### **Censoring times**

- The time of event endpoint is not observed exactly;
- Right-censoring: the censored time-to-death will be a time beyond the observed time.

$$\mathcal{L}(\boldsymbol{\theta} \mid \boldsymbol{t}) = \prod_{i=1}^{n} \left[ f(t_i; \boldsymbol{\theta}) \right]^{\mathsf{cens}_i} \left[ S(t_i; \boldsymbol{\theta}) \right]^{(1-\mathsf{cens}_i)},$$

- $t_i$  is the *i*-th recorded time, i = 1, ..., n,
- $cens_i = 0$ , if  $t_i$  is a censored time and 1 otherwise.

# <sup>3</sup> **Results and discussion**

Results and discussion

### Non-linear models (Chrysodeixis includens)



Figure: Results for *Helicoverpa armigera*. (a) fitted logistic curves and (b) parameter estimates and multiple comparisons (5% significance level).

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### Non-linear models (Helicoverpa armigera)



Figure: Results for *Helicoverpa armigera*. (a) fitted logistic curves and (b) parameter estimates and multiple comparisons (5% significance level).

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### **AFT models**

Table: Analysis of deviance for the models fitted to the time-to-death of the bollworms *Chrysodeixis includens* and *Helicoverpa armigera*.

Specie	Effect	df	Deviance	Diff df	2 logLik	p-value
Chrysodeixis	Null			187	679.9222	
2	Treatment	3	23.4077	184	656.5144	0.00003
	Experiment	1	7.7327	183	648.7817	0.00542
	Interaction	3	1.1316	180	647.6501	0.76946
Helicoverpa	Null			198	693.8224	
	Treatment	3	11.3926	195	682.4298	0.00978
	Experiment	1	1.6096	194	680.8202	0.20455
	Interaction	3	1.2429	191	679.5772	0.74272

### **AFT models**



Figure: Survival curves for the estimated times to death.

# <sup>4</sup> Final remarks

#### Data analysis results

- In this work, we evaluated the effect of fungus inoculation in soybean plants on delayed development of bollworms;
- The data analysis has shown that inoculation of fungi may delay the bollworm development.

The weight gain and time-to-death were lower for bollworms fed with fungus inoculated plants.

#### Methodological contributions

- Use of non-linear models with interpretive parameters and multiple comparison tests for each parameter;
- Use of parametric models for time-to-event data with comparison of the survival curves by using the likelihood ratio test.

#### Future research

 Joint modelling longitudinal outcomes and time-to-event data (Rizopoulos 2012).

#### Bibliography

### References

- Benjamini, Y. & Yekutieli, D. (2001), 'The control of the false discovery rate in multiple testing under dependency', *Annals of Statistics* **29**, 1165–1188.
- Bretz, F., Hothorn, T. & Westfall, P. (2016), *Multiple comparisons using R*, Chapman & Hall / CRC Press, New York.
- Kalbfleisch, J. D. & Prentice, R. L. (2002), *The statistical analysis of failure time data*, Vol. 360, 2nd edition edn, John Wiley & Sons, Hoboken, New Jersey.
- Pinheiro, J., Bates, D., DebRoy, S., Sarkar, D. & R Core Team (2018), nlme: Linear and Nonlinear Mixed Effects Models. R package version 3.1-137. URL: https://CRAN.R-project.org/package=nlme
- Pinheiro, J. C. & Bates, D. M. (2000), *Mixed-Effects Models in S and SPLUS*, Springer Series in Statistics and Computing, New York.
- Rizopoulos, D. (2012), Joint models for longitudinal and time-to-event data: With applications in R, Chapman and Hall/CRC.
- Therneau, T. M. (2015), A Package for Survival Analysis in S. version 2.38. URL: https://CRAN.R-project.org/package=survival
- Verbeke, G. & Molenberghs, G. (2000), *Linear mixed models for longitudinal data*, Springer Series in Statistics, New York.