Possibilities and Limits of Logistic Regression in a Study of the transmission dynamics of ESBL/AmpC producing E. coli between broiler flocks

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Our model world:

100 flocks, each with 100 animals

Production chain with 5 stages
### The big picture - Transmission model of ESBL/AmpC *E.coli* cont.

<table>
<thead>
<tr>
<th>parents_flock</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Prevalence in Parent FLOCKS</strong></td>
</tr>
</tbody>
</table>

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Modelling impact of horizontal transmission on flock prevalence

We saw that:

\[ \text{Prev}_0 \ (\text{initial flock prevalence}) \]

\[ \text{Prev}_1 \ (\text{flock prevalence after horizontal transmission}) \]

„Under the hood“:

Horizontal transmission:

\[ \text{Prev}_1 = \text{Prev}_0 + (100 - \text{Prev}_0) \]
The logistic regression model used

Logistic regression provides probability for binomial outcome. Here: probability that flock got colonised by ESBL/AmpC *E.coli* in given stage of the production chain.

\[
\text{Probability that flock got colonised} = \frac{1}{1 + e^{-(\text{hs} \cdot \frac{\text{prev}_0}{\text{nf}} + \text{hp} \cdot \text{col_hist} + \text{ho})}}
\]
Parameterisation of the model

Question: How to chose the values for $hs$, $hp$ and $ho$?

Answer: One fits logistic regression model to data

Challenge: We have no appropriate data (on subsequent production cycles)

Approach: Computer experiment with theoretical scenarios which means

- making our own theoretical data,
- fitting regression model to theoretical data (using R package brglm),
- get an idea of possible values ranges

Let`s look at some scenarios
Scenario 1: Intercycle transmission dominates

Intercycle transmission means: colonisation status of a house is determined by its colonisation status in previous production cycle.

and

No intrafarm transmission
Scenario 1: Intercycle transmission dominates

Fitting the logistic regression model:

<table>
<thead>
<tr>
<th></th>
<th>Value</th>
<th>P-Value (Wald test)</th>
</tr>
</thead>
<tbody>
<tr>
<td>hs</td>
<td>8.143</td>
<td>0.29047</td>
</tr>
<tr>
<td>hp</td>
<td>13.860</td>
<td>&lt; 0.001 ***</td>
</tr>
<tr>
<td>ho</td>
<td>-8.295</td>
<td>0.00357 **</td>
</tr>
</tbody>
</table>

Model of scenario 1:

\[
1 + e^{-\left(8.143 \cdot \frac{prev_0}{nf} + 13.86 \cdot \text{col_hist} - 8.295\right)}
\]
Scenario 1: Intercycle transmission dominates

520 predictions, using a threshold of 0.5
the model identifies 390 true negatives and
130 true positives

⇒ Accuracy = \[
\frac{520}{520} = 100%
\]
Intercycle transmission means: colonisation status of a house correlates highly correlated with its colonisation status in the previous production cycle.

When a house was colonised in previous cycle it has 90% probability to get colonised in the current cycle.
Scenario 2: Intercycle transmission dominates - probabilistic

Fitting the logistic regression model:

<table>
<thead>
<tr>
<th></th>
<th>Value</th>
<th>P-Value (Wald test)</th>
</tr>
</thead>
<tbody>
<tr>
<td>hs</td>
<td>-1.0632</td>
<td>0.309</td>
</tr>
<tr>
<td>hp</td>
<td>5.2550</td>
<td>&lt; 0.001 ***</td>
</tr>
<tr>
<td>ho</td>
<td>-2.6993</td>
<td>&lt; 0.001 ***</td>
</tr>
</tbody>
</table>

520 predictions, using a threshold of 0.5, the model identifies 349 true negatives and 140 true positives.

⇒ Accuracy = \( \frac{489}{520} \approx 94\% \)
Scenario 3: No intrafarm and No intercycle transmission

Each Farm has exactly one colonised house

And

House was never colonised in previous production cycle
Scenario 3: No intrafarm and No intercycle transmission

Fitting the logistic regression model:

<table>
<thead>
<tr>
<th></th>
<th>Value</th>
<th>P-Value (Wald test)</th>
</tr>
</thead>
<tbody>
<tr>
<td>$hs$</td>
<td>-73.893</td>
<td>$&lt; 0.001$ ***</td>
</tr>
<tr>
<td>$hp$</td>
<td>1.550</td>
<td>0.448</td>
</tr>
<tr>
<td>$ho$</td>
<td>5.559</td>
<td>$&lt; 0.001$ ***</td>
</tr>
</tbody>
</table>

520 predictions, using a **threshold of 0.5**
the model identifies 390 true negatives and 130 true positives

⇒ Accuracy = \[
\frac{520}{520} = 100\% \]
Scenario 4: Intrafarm but no explicit intercycle transmission

On each farm all houses except one is colonised
Scenario 4: Intrafarm but no explicit intercycle transmission

Fitting the logistic regression model:

<table>
<thead>
<tr>
<th></th>
<th>Value</th>
<th>P-Value (Wald test)</th>
</tr>
</thead>
<tbody>
<tr>
<td>( hs )</td>
<td>-11.0713</td>
<td>&lt; 0.001 ***</td>
</tr>
<tr>
<td>( hp )</td>
<td>0.9272</td>
<td>0.00102 **</td>
</tr>
<tr>
<td>( ho )</td>
<td>7.4618</td>
<td>&lt; 0.001 ***</td>
</tr>
</tbody>
</table>

520 predictions, using a threshold of 0.6 the model identifies 86 true negatives and 367 true positives

\[ \Rightarrow \text{Accuracy} = \frac{453}{520} \approx 87\% \]
Scenario 5: Random colonisation

Each house has a 50% chance of becoming colonised
Scenario 5: Random colonisation

Fitting the logistic regression model:

<table>
<thead>
<tr>
<th></th>
<th>Value</th>
<th>P-Value (Wald test)</th>
</tr>
</thead>
<tbody>
<tr>
<td>hs</td>
<td>0.005857</td>
<td>0.989</td>
</tr>
<tr>
<td>hp</td>
<td>0.16887</td>
<td>0.336</td>
</tr>
<tr>
<td>ho</td>
<td>-0.132886</td>
<td>0.499</td>
</tr>
</tbody>
</table>

520 predictions, using a **threshold of 0.5**
the model identifies 138 true negatives and 133 true positives

⇒ **Accuracy** = \( \frac{271}{520} \approx 52\% \)
Summarising Results

Based on the five scenarios value ranges for the coefficients $hs, hp$ and $ho$ were found and used in the transmission model.

<table>
<thead>
<tr>
<th></th>
<th>Minimum</th>
<th>Maximum</th>
</tr>
</thead>
<tbody>
<tr>
<td>$hs$</td>
<td>-73.89</td>
<td>8.14</td>
</tr>
<tr>
<td>$hp$</td>
<td>0.17</td>
<td>13.86</td>
</tr>
<tr>
<td>$ho$</td>
<td>-8.30</td>
<td>7.46</td>
</tr>
</tbody>
</table>
Using the parameterisation in broiler production chain model
Conclusion Possibilities and limits

Possibilities:
Logistic model useful in predicting outcomes
Helps identify directions which to investigate further

Limits:
Numeric values of regression coefficients have no direct real world interpretation
At the current stage our regression model is limited in grain size of analysis and therefore limited in giving particular hints on interventions for farmers
Thank you for your attention

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Backup
Choosing thresholds – ROC curve

Receiver Operating Characteristic curve = ROC curve for scenario 4

AUC = 0.82
Predicting with the model

Prediction via threshold:

Example – for given independent variables and coefficients say that

\[
\text{Probability that flock got colonised} = \frac{1}{1 + e^{-(2.211 \cdot 0.5 + 1 \cdot 0 - 0.7)}} \approx 0.6
\]

Introduce threshold, say 0.5

Prediction: Flock is predicted to become colonized if probability > threshold

In our case: since 0.6 > 0.5 the flock is predicted to become colonized
How accurate are model predictions?

Accuracy = number of correct prediction / number of predictions

Example: Introduce threshold, say 0.5

Probability that flock got colonised = \[
\frac{1}{1 + e^{-\left(8.143 \cdot \frac{\text{prev}_0}{n_f} + 13.86 \cdot \text{col}_\text{hist} - 8.295\right)}}
\]

<table>
<thead>
<tr>
<th>Farm with 3 broiler houses</th>
<th>Prediction for Production Cycle 1</th>
<th>Prediction for Production Cycle 2</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>0.9997</td>
<td>0.0538</td>
</tr>
<tr>
<td></td>
<td>0.9997</td>
<td>0.0038</td>
</tr>
</tbody>
</table>

Accuracy = \(\frac{1}{3} \approx 33\%\)