Modeling codling moth (Lepidoptera: Tortricidae) phenology and predicting egg hatch in apple orchards of the Maule Region, Chile

Wilson Barros-Parada1, Alan L. Knight2, and Eduardo Fuentes-Contreras1*

Studies were conducted in the Maule Region to characterize the phenology of the codling moth, Cydia pomonella (L.). The Predictive Extension Timing Estimator (PETE) and a logistic phenological model were validated with eight data sets of cumulative moth catches in sex pheromone (PH) and kairomone-baited traps and the cumulative occurrence of fruit injuries from apple (Malus domestica Borkh.) orchards during the 2009-2011 seasons. Second, the start of egg hatch was predicted from the first sustained male and female moth catches (biofix) in traps baited with pear ester (PE), PE+acetic acid (AA), PE+PH, and PH alone. Both phenological models fit data well except that the logistic provided a better fit than the PETE model of the phenomenology of egg hatch of the codling moth in the first generation, with a difference of 11 d between models in the prediction of 50% egg hatch. No significant difference was found between biofix dates established for males using either PH or PE+PH lures or for the biofix date based on female catches with PE+AA or PH+PE. The biofix established with the sustained female catch occurred nearly 11 d later than the male-based biofix. The use of a female biofix provided on average a 4-d improvement in the prediction of first egg hatch compared with the traditional use of a male biofix, but this difference was not significant. The use of PE+AA lures increased the proportion of cases when a female-based biofix could be established compared with the use of the PH+PE lure.

Key words: Biofix, Cydia pomonella, degree days, codlemone, Malus domestica, pear ester.

INTRODUCTION

Phenological models have been important tools used to improve the management of codling moth, Cydia pomonella (L.) (Lepidoptera: Tortricidae) in many apple producing areas worldwide (Tomkins et al., 1987; Knight and Croft, 1991; Blago, 1992; Ahmad et al., 1995; Boivin et al., 2005; Samietz et al., 2007; Knight, 2007; Jones et al., 2008; 2010; 2013). The Predictive Extension Timing Estimator (PETE) model was the first widely-adopted phenological model developed for codling moth in the USA, and is based on a time-varying distributed delay algorithm (Welch et al., 1978). The validation of PETE codling moth model was performed successfully with data of accumulated male adult catches in sex pheromone-traps (PH) and egg-hatching in relation to degree days (DD) (Riedl et al., 1976). Subsequently, PETE models were implemented in many apple producing states in the USA (Jorgensen et al., 1979; Welch et al., 1981; Brunner et al., 1982; Croft and Knight, 1983; Pitcairn et al., 1992; Beers and Brunner, 1992). Efforts to validate PETE on apple pests in Chile started in the 1980’s (Croft and Knight, 1983; Vial, 1987), but growers’ interest in its implementation was low due to their reliance on repeated calendar-based insecticide sprays in order to meet the strong quarantine restrictions imposed by export markets (González, 2003). Currently, the PETE codling moth model is available in Chile on a web-based service (http://www.agroclima.cl) to farmers associated with “Fundación para el Desarrollo Frutícola” (FDF).

A new phenology model was developed for codling moth in Washington State (USA), because of the reported poor fit of the PETE model that many growers observed in this region once sex pheromones began to be adopted for mating disruption and insecticide spray programs became more limited (Knight, 2007). The cumulative curves for both adult emergence and egg hatch based on male catches in sex pheromone-traps and newly injured fruits were fit to logistic equations for both generations of codling moth, and the new model predicted a delay of 100 degree-days (DD) from the PETE model for 50% egg hatch in the first generation (Knight, 2007). The potential impact of this difference between models on the effectiveness of insecticide programs targeting eggs and neonate larvae of the codling moth was shown conceptually to be an important concern for effective insecticide-based management of this pest. This concern is heightened today as growers must rely on the use of selective insecticides which often have shorter residual activity than the previous broad spectrum insecticides.
Identification of these associated negative impacts. Growers to further reduce their use of insecticides and key factors affecting fruit production. Development of a recognition by apple growers in Chile that the failures (Knight, 2007).

Need to be accurate at a temporal scale consistent with phenology models available to aid management decisions (Brunner et al., 2005; Jones et al., 2010). Therefore, development and implementation of accurate phenology models for codling moth is important has increased (González, 2003). Here, apple production is primarily export-driven and due to severe quarantine restrictions for codling moth-injured fruits growers’ reliance on insecticide use is high. Efforts to minimize residues on fruit and improve the safety of workers and avoid disturbing the population dynamics of secondary pests, such as San José scale, Diaspidiotus perniciosus (Comstock), and woolly apple aphid, Eriosophia lanigerum (Hausmann), are key factors affecting fruit production. Development of a more accurate predictive phenology model should allow growers to further reduce their use of insecticides and these associated negative impacts.

Identification of (E, Z)-2,4-ethyl-decadienoate (pear ester, PE) as a potent attractant for both sexes of codling moth (Light et al., 2001), as well as the use of acetic acid (AA) as a synergist to increase the catch of both sexes (Landolt et al., 2007), has allowed the development of new traps and lures for coding moth monitoring (Knight, 2010a; 2010b; 2010c; Barros-Parada et al., 2013). These new technologies provide potential alternatives for the definition of more accurate biological events to start the accumulation of DD in phenological models (Knight and Light, 2005a; 2005b).

Biofix was originally defined as “the first male moth or moths in the traps baited with the sex pheromone, (E,E)-8,10-dodecadien-1-ol (PH), with no significant interruption in catches thereafter” (Riedl et al., 1976). This definition of a sustained biofix has been further adjusted with the occurrence of suitable dusk temperatures for moth sexual activity (> 15.6 °C) (Pickel et al., 1986; Blago, 1992) or suitable maximum daily temperatures (> 21.1 °C) (Pitcairn et al., 1990). More recently, Knight and Light (2005a) using traps baited with PE re-defined the biofix for male, female or any sustained catches with an adjustment for maximum daily temperature.

Herein, we report the results from studies aimed to i) statistically validate the codling moth PETE (Brunner et al., 1982) and logistic models (Knight, 2007) for adult male catches and egg hatch in the Maule Region of central Chile; and ii) compare the establishment of a biofix based on sustained coding moth male and/or female catches in traps baited with PE+AA, PE+PH, or PH.

MATERIALS AND METHODS

Field studies
Five abandoned apple (Malus domestica Borkh.) orchards in the Maule Region, Chile, were monitored during the 2009-2010 and 2010-2011 seasons. The Pencahue site (35°23'09” S, 71°48'39” W) was a 1.3 ha ‘Red King Oregon’, ‘Royal Gala’, and ‘Fuji’ orchard; the Colín orchard (35°27'56” S, 71°44'04” W) was a 7.8 ha mixed block of ‘Red Chief’, ‘Royal Gala’ and ‘Granny Smith’, and the Talca site (35°27'26” S, 71°36'33” W) was a 1-ha mixed block of ‘Fuji’ and ‘Royal Gala’. All of these three sites were monitored in both seasons. The Villa Alegre orchard (35°01'12” S, 71°14'26” W) was a 5.8 ha mixed block of ‘Granny Smith’ and ‘Royal Gala’ monitored only in 2009-2010. The Rauco orchard (34°55'57.58” S, 71°16'38” W) was a 0.5 ha mixed block of ‘Red King Oregon’, ‘Granny Smith’, and ‘Royal Gala’ monitored only in 2010-2011. The mean tree height in orchards varied from 1.8 to 5.4 m, and the tree densities were between 666-1250 trees ha⁻¹. All orchards had been abandoned for > 8 yr, and no insecticide sprays or mating disruption technologies were used in any of the sites during the 2-yr study.

Trees (3-20 per site) with a full fruit load were randomly selected in each orchard at the beginning of the season following bloom to provide an estimated population of 2000-3500 fruits per site. During both seasons, trees selected at each site were sampled once a week. Two technicians examined all fruits on designated trees using ladders on each date. Sampling efforts on each date ranged from 1 to 2 h per orchard depending on the number of fruits checked, number of fruits removed, and tree size. Degree days (DD) were calculated with a simple sine wave function using 10 °C and 31.1 °C as temperature thresholds with a horizontal cutoff (Baskerville and Emin, 1969). The accumulated mean DD (SE) between sampling dates was 59.2 (1.3) for the first generation and 95.4 (3.0) for the second generation of codling moth. A variable proportion of damaged fruit evaluated in all orchards had multiple injuries. Data on the total number of injuries were recorded, instead of the number of injured fruits. On each sampling date damaged fruits were removed to avoid counting them on the next evaluation date. The proportion of fruits left on sampled trees, after removal of fruits with injuries, by the end of the season was 0.05-0.8 in all sites, except for the first and second generation at Colín during 2009-2010, where little damage occurred and these data were not included in the study. Also, in the Pencahue orchard during 2009-2010 the proportion of fruit injury was > 0.95 by mid-season, and data for the second generation were not included in the analyses.

Male and female codling moth were monitored in each orchard with five trap-lure combinations during the 2009-2010 season: i) clear delta-pear ester (PE) (Pherocon DA, Trécé Inc., Adair, Oklahoma, USA) + acetic acid (AA); ii) clear round-PE+AA, iii) orange delta-PE+ sex pheromone (PH) (Pherocon CM-DA Combo, Trécé Inc., Salinas, California, USA); iv) white delta-PE+ PH, and v) white delta-PH (Pherocon CM L2, Trécé Inc., Salinas). Clear delta and round delta-PE+AA as well as orange delta and white traps were baited with the sex pheromone, (E,E)-8,10-dodecadien-1-ol (PH), with no significant interruption in catches thereafter” (Riedl et al., 1976). This definition of a sustained biofix has been further adjusted with the occurrence of suitable dusk temperatures for moth sexual activity (> 15.6 °C) (Pickel et al., 1986; Blago, 1992) or suitable maximum daily temperatures (> 21.1 °C) (Pitcairn et al., 1990). More recently, Knight and Light (2005a) using traps baited with PE re-defined the biofix for male, female or any sustained catches with an adjustment for maximum daily temperature.

Herein, we report the results from studies aimed to i) statistically validate the codling moth PETE (Brunner et al., 1982) and logistic models (Knight, 2007) for adult male catches and egg hatch in the Maule Region of central Chile; and ii) compare the establishment of a biofix based on sustained coding moth male and/or female catches in traps baited with PE+AA, PE+PH, or PH.
delta-PE+PH showed similar catches of codling moth, and therefore only one of each was used in the following season. During the 2010-2011 season three trap-lure combinations were used: i) clear delta-PE+AA, ii) orange delta-PE+PH, and iii) white delta-PE. The AA lure was prepared with 6 mL glacial acetic acid (99%) (Sigma-Aldrich, St. Louis, Missouri, USA) added to a 8.0 mL polyethylene vial (Nalg-Nunc International, Rochester, New York, USA) with a 3.0 mm hole in the screw cap. A 10 g cotton ball was put into each vial as described by Barros-Parada et al. (2013). Clear delta and round traps were made of plastic film as described by Knight (2010b) and Barros-Parada et al. (2013). Traps were attached to poles in the upper third of the tree canopy (approx. 3 m high) according to the protocol described by Knight et al. (2006). An array of traps was placed with an N-S orientation with traps spaced 30-50 m apart and ≥ 10 m from the edge of the orchard. Lures were replaced every 8 wk and the sticky liners when needed. Traps were checked two or three times per week and rotated within the array on each sampling date. Moths were removed from traps on each date and sexed in the field (Riedl et al., 1976). Daily temperature was recorded with a data-logger (Hobo, Bourne, Massachusetts, USA) located in the tree canopy at each sampling site. Total male trap catches and egg hatch for each flight and season were compared with repeated measures ANOVA.

Validation of phenological models

Accumulated male moth catches in sex pheromone-baited traps as a function of DD accumulations during the season were used to calculate the proportion of the flight cycle for each generation of codling moth. Since no significant differences in the male accumulated proportion of catches between trap-lure combinations (PH and PE+PH) were found, the pooled catches were used to evaluate the fit of the models.

The biofix in each orchard-year was established from sustained male moth catch. The moth catch that occurred on the biofix date was assigned the total DD accumulated on the day the trap was checked. The fruit damage was used as an estimation of egg hatch, assuming the occurrence of fruit injury 1 d after egg hatching as suggested by Knight (2007). Therefore, registered DD for each orchard in the day before the detection of fruit damage was also accumulated. The duration of codling moth flight and the egg hatch period for the first and second generation were set as 444 and 567 and 1044 and 1189 DD after biofix, respectively (Brunner et al., 1982). This approach was used because it is not possible to clearly distinguish between the first and second generations of codling moth due to some overlapping development.

The statistical validation of these data fit to the logistic model (WA logistic) developed for Washington by Knight (2007) and a logistic version of the PETE model (Brunner et al., 1982) was performed. The PETE logistic model was developed from the tabular data of the cumulative moth flight and egg hatch as a function of DD available in Brunner et al. (1982). The fit between observed and predicted values was compared with the coefficient of determination ($R^2$), root mean square error (RMSE), and mean absolute error (MAE) defined by Mayer and Butler (1993). Finally, the differences in DD between model predictions for 10%, 50%, and 90% completion of adult male flight and egg hatch were calculated.

Biofix determination

Several types of trap catch data from each orchard were used to establish a biofix for male and female adult emergence: i) sustained male catch in traps baited with PH, PE+AA or PE+PH lures; and ii) sustained female catch in traps baited with PE+PH or PE+AA lures. Data from different types of traps were pooled in these determinations. A strict definition of a biofix was adhered to as the date of sustained catch in which at least two consecutive days with maximum temperatures above 21.1 °C occurred between trap checks, with the following modifications from the definition described by Knight and Light (2005a). First, sustained catch in this study was defined as the date when two consecutive trap checks caught at least one individual moth, instead of an average catch of one moth per trap. Second, sustained catch was established at the start of the period of the second consecutive individual catch, instead of the start of the period from the first of the consecutive catches above one moth per trap. The Julian date (JD) of these defined biofix’s was used to compare trap-lure treatments. The cumulated DD from the observed to the expected occurrence of egg hatch were estimated using the backtracking data from Knight and Light (2005a). Observed egg hatch was obtained from fruit damage evaluations, based on the assumption that egg hatch started 1 d after the previous sampling date without damage detection. The mean and (SE) DD between fruit damage detection and observed egg hatch was 29.6 (4.4) DD. Differences between biofix as JD, and in the observed and expected dates and DD totals for egg hatch were analyzed with a non-parametric Kruskal-Wallis test.

RESULTS

Seasonal male catches and egg hatch

The cumulative catches of codling moth males in the traps were not significantly different between generations ($F_{1,6} = 0.04, P = 0.85$), seasons ($F_{1,6} = 1.94, P = 0.21$) or with the interaction of these two factors ($F_{1,6} = 0.83, P = 0.40$). The numbers of fruit injuries was significantly lower in the first than the second year ($F_{1,4} = 9.93, P < 0.05$), but not between generations ($F_{1,4} = 1.43, P = 0.30, NS$) nor was the interaction Year × Generation significant ($F_{1,4} = 3.07, P = 0.15$).

Model validation

The logistic and the PETE logistic fit models showed similar $R^2$, MAE, and RMSE for both adult male
emergence periods (Table 1). However, the WA logistic model for the first generation egg hatch had a higher $R^2$ and lower MAE and RMSE than those obtained for the PETE logistic model (Table 1). In contrast, both models had similar $R^2$, MAE, and RMSE values for the second generation egg hatch period (Table 1).

The difference between the WA logistic and the PETE logistic model of adult male emergence was 22 DD and -4 DD for 50% of the first and second flight, which correspond approximately to 3 d and less than 1 d for the first and second flight, respectively (Table 2). On the contrary, for the first egg hatch period the 50% egg hatch showed a difference of 81 DD between models (Table 2). This difference was only 7 DD for the second egg hatch period (Table 2). Since mean orchard temperatures were lower in spring than in summer, the difference between models in DD has a stronger effect in calendar days for the first (11 d) than for the second (1 d) egg hatch period. Such differences resulted in a more accurate prediction of the first egg-hatch by the WA logistic than by PETE logistic model, which showed -50 DD and -130 DD difference with the mean egg-hatch detected in the field, respectively (Table 2).

### Table 1. Values of statistical validation criteria used to compare the WA logistic and the PETE logistic model for codling moth phenology during the first and generation in apple orchards during 2009-2011 from the Maule Region, Chile.

<table>
<thead>
<tr>
<th>Event</th>
<th>Model</th>
<th>Number of data points</th>
<th>$R^2$</th>
<th>RMSE</th>
<th>MAE</th>
</tr>
</thead>
<tbody>
<tr>
<td>1st Flight</td>
<td>WA-Logistic</td>
<td>571</td>
<td>0.79</td>
<td>0.21</td>
<td>0.15</td>
</tr>
<tr>
<td></td>
<td>PETE-Logistic</td>
<td>571</td>
<td>0.78</td>
<td>0.19</td>
<td>0.13</td>
</tr>
<tr>
<td>2nd Flight</td>
<td>WA-Logistic</td>
<td>438</td>
<td>0.78</td>
<td>0.18</td>
<td>0.13</td>
</tr>
<tr>
<td></td>
<td>PETE-Logistic</td>
<td>438</td>
<td>0.78</td>
<td>0.18</td>
<td>0.13</td>
</tr>
<tr>
<td>1st Egg hatch</td>
<td>WA-Logistic</td>
<td>114</td>
<td>0.94</td>
<td>0.11</td>
<td>0.06</td>
</tr>
<tr>
<td></td>
<td>PETE-Logistic</td>
<td>114</td>
<td>0.78</td>
<td>0.27</td>
<td>0.18</td>
</tr>
<tr>
<td>2nd Egg hatch</td>
<td>WA-Logistic</td>
<td>69</td>
<td>0.90</td>
<td>0.11</td>
<td>0.09</td>
</tr>
<tr>
<td></td>
<td>PETE-Logistic</td>
<td>69</td>
<td>0.89</td>
<td>0.13</td>
<td>0.09</td>
</tr>
</tbody>
</table>

$R^2$: Coefficient of determination, higher values denote a better data fit. All regressions were significant $P < 0.05$; RMSE: root mean square error, lower values denote a better fit of data; MAE: mean absolute error, lower values denote a better fit of data.

### Table 2. Observed and predicted degree-days (DD) values of the WA logistic and the PETE logistic models for codling moth flight and egg hatch in apple orchards during 2009-2011 seasons from the Maule Region, Chile.

<table>
<thead>
<tr>
<th>Event (% completion)</th>
<th>DD value for event</th>
<th>WA-Logistic and field data</th>
<th>PETE-Logistic and field data</th>
<th>Logistic models</th>
</tr>
</thead>
<tbody>
<tr>
<td>1st Flight</td>
<td>10</td>
<td>54 (28)</td>
<td>93</td>
<td>39</td>
</tr>
<tr>
<td>50</td>
<td>164 (72)</td>
<td>170</td>
<td>148</td>
<td>6</td>
</tr>
<tr>
<td>90</td>
<td>362 (50)</td>
<td>289</td>
<td>271</td>
<td>-73</td>
</tr>
<tr>
<td>2nd Flight</td>
<td>10</td>
<td>621 (103)</td>
<td>595</td>
<td>-26</td>
</tr>
<tr>
<td>50</td>
<td>812 (74)</td>
<td>748</td>
<td>752</td>
<td>-64</td>
</tr>
<tr>
<td>90</td>
<td>975 (48)</td>
<td>901</td>
<td>911</td>
<td>-74</td>
</tr>
<tr>
<td>1st Egg hatch</td>
<td>10</td>
<td>293 (36)</td>
<td>257</td>
<td>-36</td>
</tr>
<tr>
<td>50</td>
<td>413 (28)</td>
<td>364</td>
<td>283</td>
<td>-49</td>
</tr>
<tr>
<td>90</td>
<td>520 (51)</td>
<td>472</td>
<td>393</td>
<td>-48</td>
</tr>
<tr>
<td>2nd Egg hatch</td>
<td>10</td>
<td>677 (53)</td>
<td>709</td>
<td>-36</td>
</tr>
<tr>
<td>50</td>
<td>966 (81)</td>
<td>892</td>
<td>885</td>
<td>-74</td>
</tr>
<tr>
<td>90</td>
<td>1120 (56)</td>
<td>1074</td>
<td>1045</td>
<td>-46</td>
</tr>
</tbody>
</table>

### DISCUSSION

Codling moth management in Chile is characterized as an intensive program of repeated insecticide sprays timed to maintain effective residues which protect the fruit from larval injuries (González, 2003). Failure to protect the fruit, even with a low proportion of larval infestation, produces the quarantine rejection of the pome fruit exports to the most profitable markets (e.g. Taiwan). When growers rely primarily on a number of broad spectrum insecticides with long residual effects (e.g. organophosphate, OP) the accuracy and precision of codling moth phenological predictions were not a major concern. However, with the replacement of OP with selective and shorter residual effect insecticides (Brunner et al., 2005; Jones et al., 2010) the improvements of phenological estimations for codling moth control are more relevant (Knight, 2007).

Our work found that the logistic model developed by Knight (2007) and the widely used PETE model (Brunner et al., 1982) were both able to predict the phenology of codling moth male flight in Maule Region, Chile. However, the logistic model appeared to predict 50% egg hatch for the first codling moth generation better than the PETE model, with a difference of 80 DD or about 11 d between
Furthermore, PE+AA lures seem to be more reliable than lures occurred significantly later than a male-based biofix. However, we found that a female-based biofix using PE+AA or PE+PH lures occurred significantly later than a male-based biofix. The use of a female-based biofix showed a trend to predict more closely the start of egg hatch in either JD or DD units compared with the standard sustained male catch in PH or PE+PH traps, but this difference was not large enough to be significant in our study. On the contrary, this difference was significant in an earlier study for males (Knight and Light, 2005a). Likely, the wide sampling interval used in or study (60 DD in spring and 95 DD in summer) limited our ability to more closely differentiate the timing of first egg hatch in relation with shorter sampling intervals achieved by Knight (2007) (36.6-83.0 DD).

**CONCLUSIONS**

The Predictive Extension Timing Estimator (PETE) and logistic models for codling moth phenology adequately estimated the adult male flight periods of the codling moth in the Maule Region, Chile. However, the logistic model showed a better estimation of the first egg-hatch period than the PETE model. This difference between models can reach up to 10 d and could produce a mismatch between the first insecticide spray and egg-hatch of neonate larvae that can damage the fruits. Biofix from sustained catch of females with pear ester and sex pheromone (PE+PH) and (PE+ acetic acid [AA]) lures was significantly delayed in relation with the standard biofix obtained from sustained catch of males in PH or PE+PH lures. PE+AA baited traps were more reliable than PE+PH to establish a female-based biofix due to their higher propensity to catch females (Knight, 2010a; 2010b; 2010c; Barros-Parada et al., 2013). Because, the male biofix estimated with PE+AA in relation with PE+PH and PH lures was not significantly different, the PE+AA lure can be used to estimate both male and female biofix. The use of a female-based biofix showed a trend to predict more closely the start of egg hatch in either JD or DD units compared with the standard sustained male catch in PH or PE+PH lures. The use of female-based biofixes proved the trend to more closely the start of egg hatch in either JD or DD units compared with the standard sustained male catch in PH or PE+PH lures.

**ACKNOWLEDGEMENTS**

This work was funded by FONDECYT grant 1071029 to EFC, and Millenium Nucleus (Molecular Ecology and Evolutionary Applications in the Agroecosystem) grant NC 120027.

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Table 3. Mean (SE) Julian date of male or female codling moth biofix established with pear ester plus acetic acid (PE+AA), pear ester plus sex pheromone (PE+PH), and sex pheromone (PH) lures, and the differences expressed as Julian dates and degree-days (DD) between predicted and observed first egg hatch in apple orchards during 2009-2011 seasons from Maule Region, Chile.

<table>
<thead>
<tr>
<th>Lure</th>
<th>Julian date of biofix*</th>
<th>Difference between observed and expected egg hatch</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Maleb</td>
<td>Femaleb</td>
</tr>
<tr>
<td>PE+AA</td>
<td>284.8 (3.9) a</td>
<td>285.3 (2.7) a</td>
</tr>
<tr>
<td>PH+PE</td>
<td>275.5 (2.3) b</td>
<td>287.3 (2.2) a</td>
</tr>
<tr>
<td>PH</td>
<td>274.8 (1.9) b</td>
<td>-</td>
</tr>
</tbody>
</table>

*Biofix was defined as the date when two consecutive checks caught at least one moth, and the maximum daily temperature was higher than 21.1 °C on at least two consecutive days from the previous trap check date.

**}
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