



Assessing the mismatch between incubation and latency for a vector-borne plant disease

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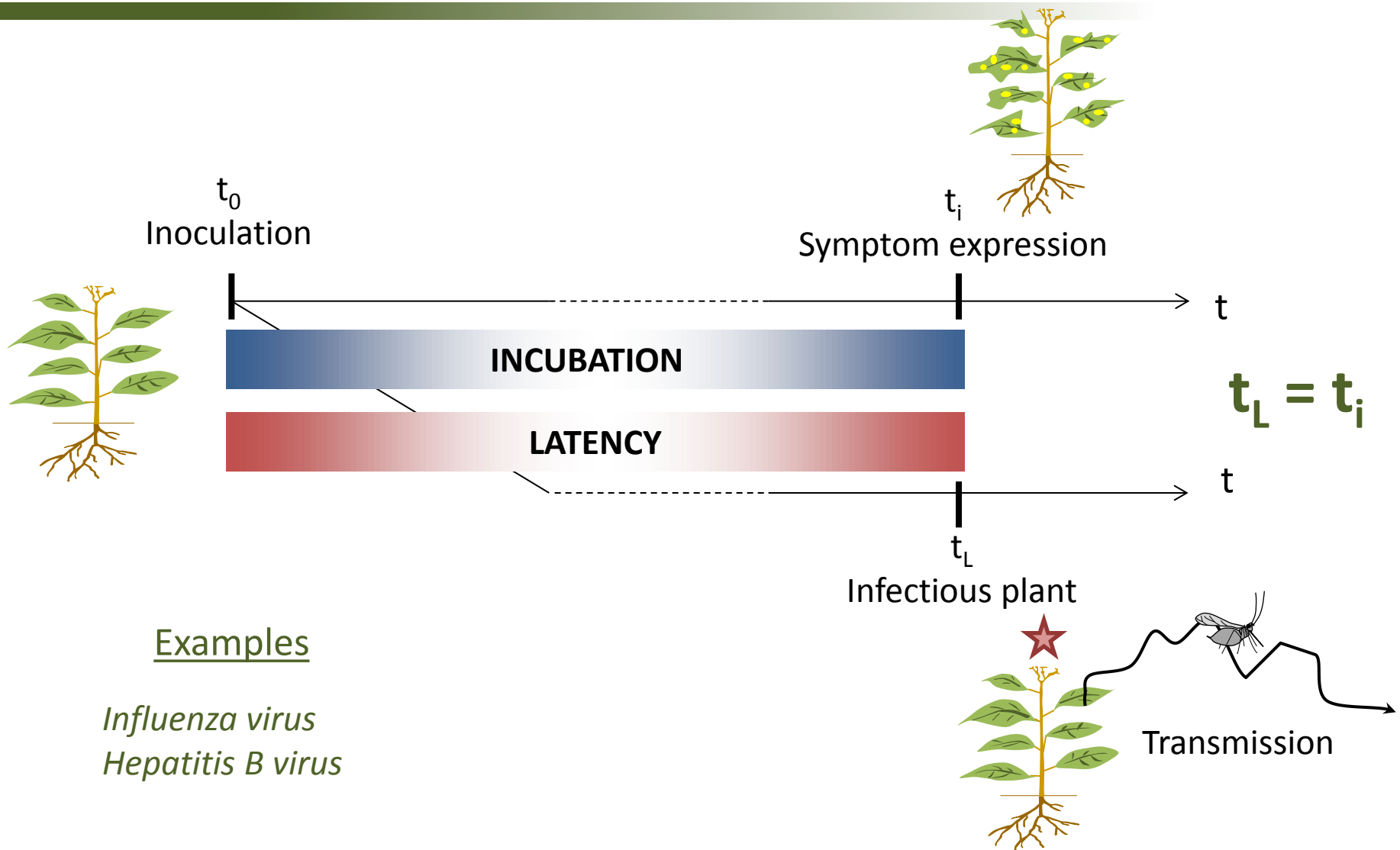


Assessing the mismatch between incubation and latency for a vector-borne plant disease

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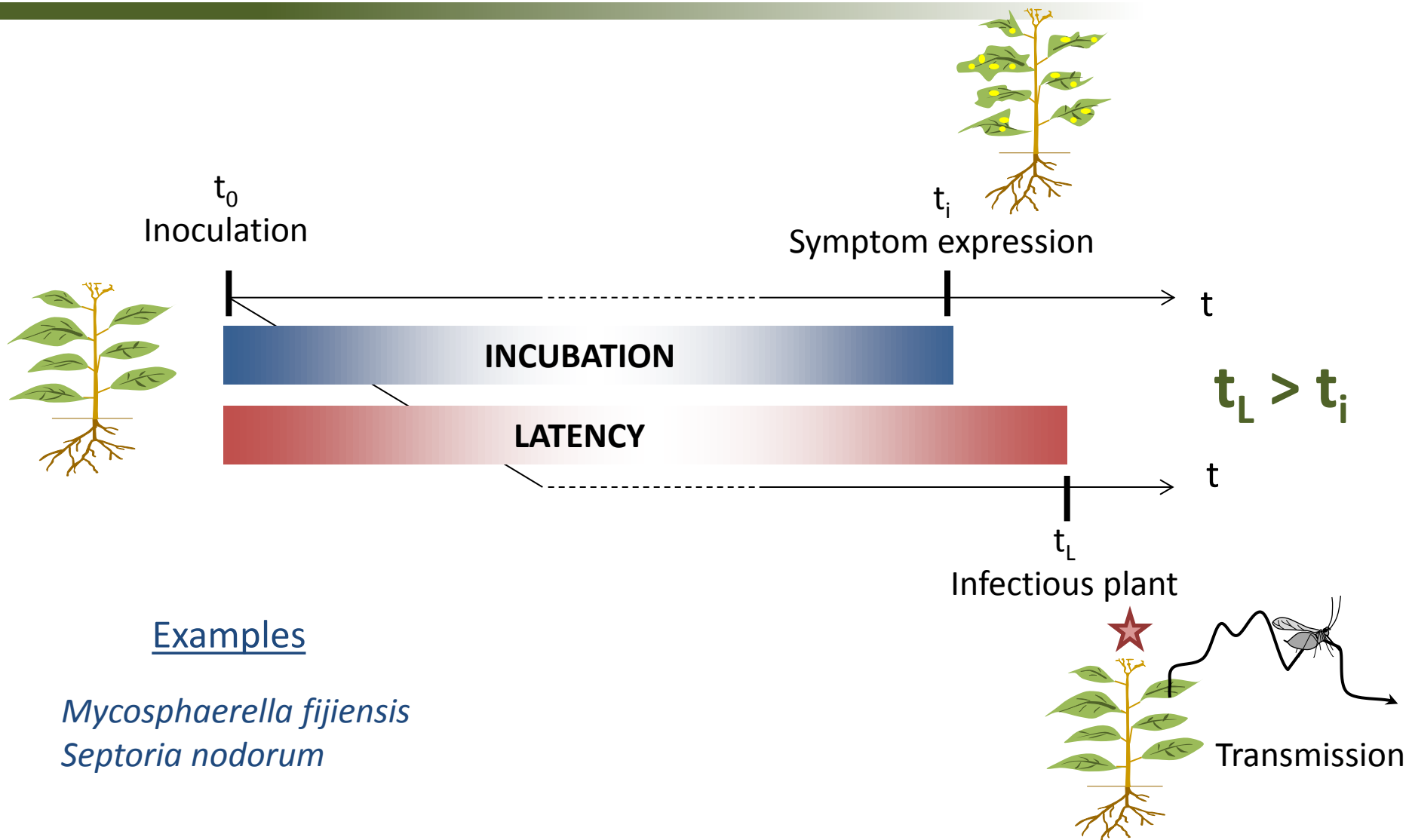
Latency vs. Incubation



Examples

Influenza virus
Hepatitis B virus

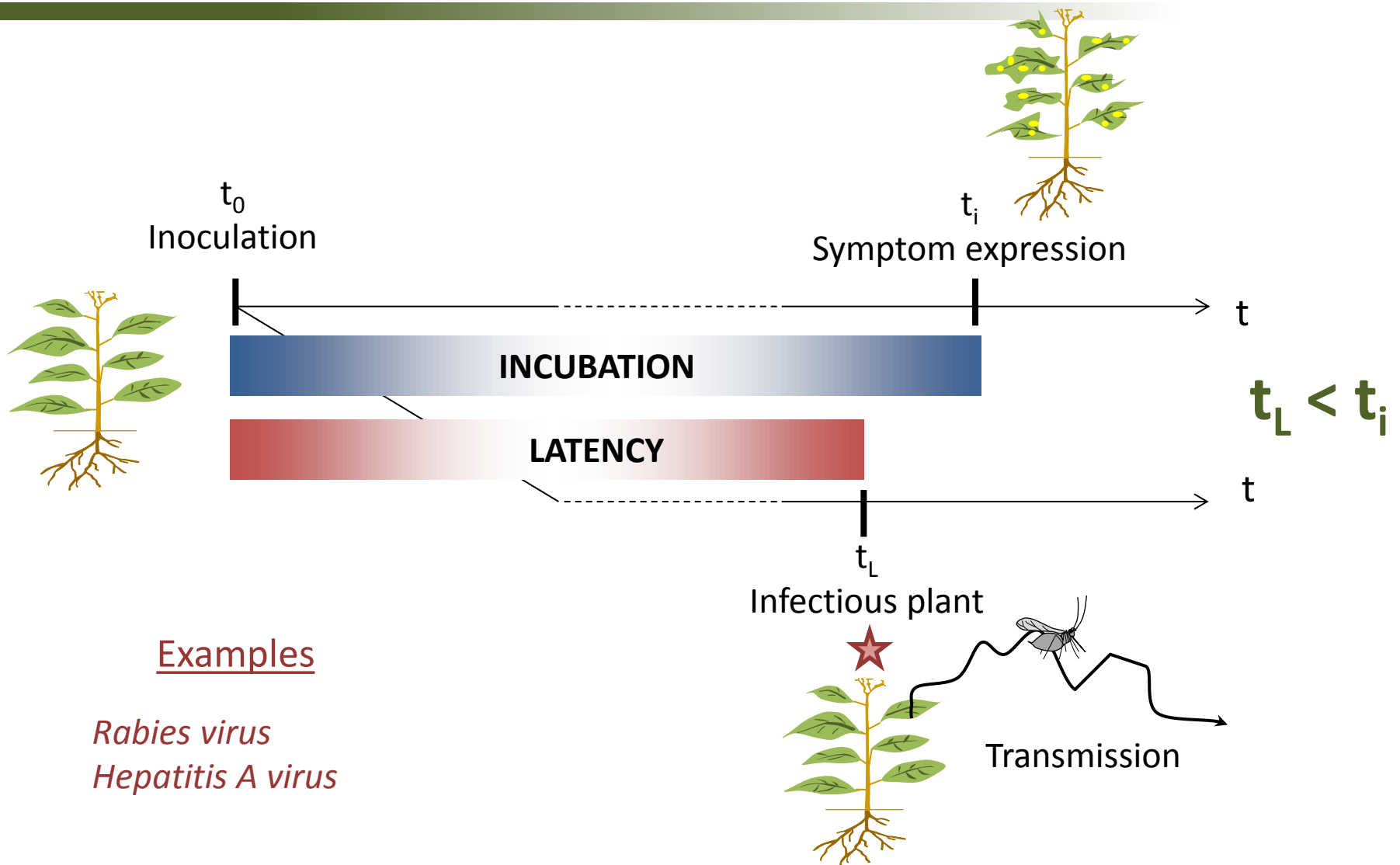
Latency vs. Incubation



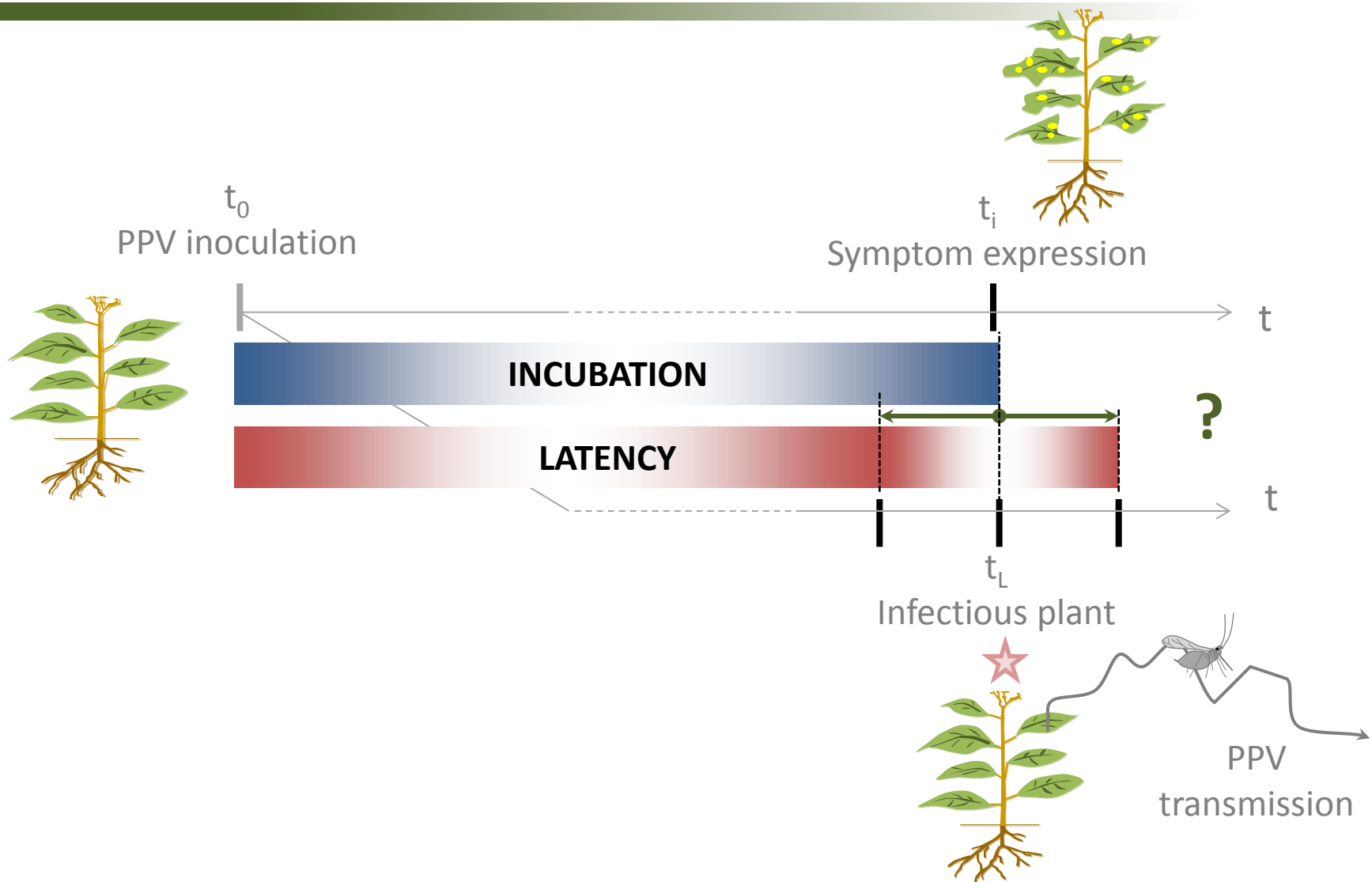
Examples

Mycosphaerella fijiensis
Septoria nodorum

Latency vs. Incubation



Latency vs. Incubation



Plum pox virus, the causal agent of sharka disease

Disease	Sharka Most damaging disease on <i>Prunus</i> 10 billions Euros of economic losses worldwide in 30 years
Pathogen	<i>Plum pox virus (PPV)</i> <i>Potyvirus</i>
Vectors	Aphids: > 20 species Human: transfer of infected material
Hosts of economic interest	<i>Prunus</i> e.g.: apricot, plum and peach trees



Sharka management strategy in France

Since the 1990's

- Frequent visual inspections of the orchards
- Removal of the symptomatic trees (or whole orchards)
- Protection of the nurseries

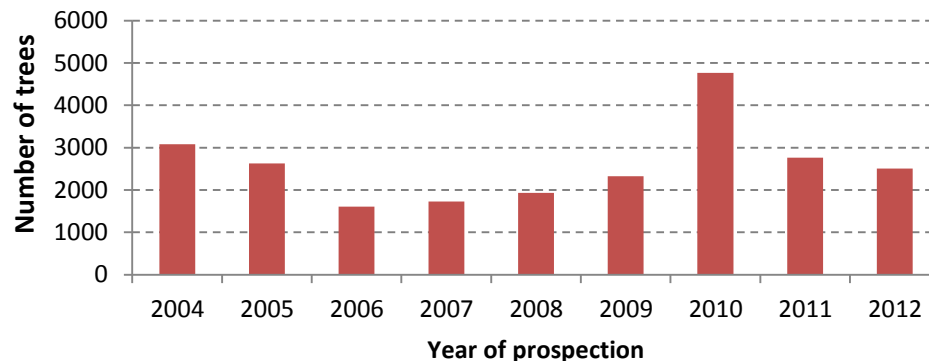
**Infected trees without symptoms
cannot be detected?**



Disappointing outcomes:

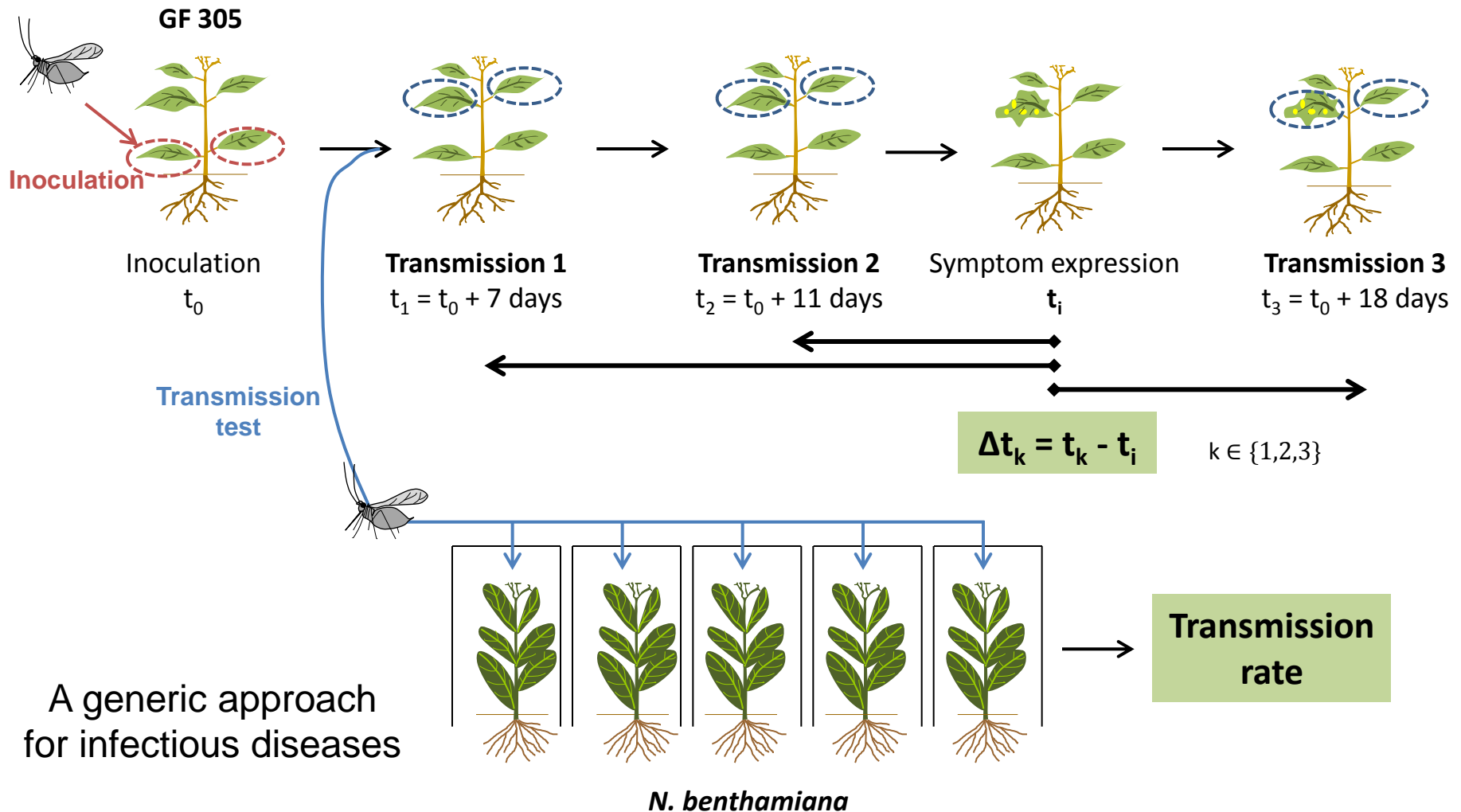
- Costly strategy
- Still many trees infected each year

Symptomatic trees detected in Gard (South of France)



(Data from regional offices of French food agency)

Experimental approach



Symptom monitoring

Definition of 5 classes of leaves:



0
No
symptom



1
Light
symptoms



2
Intermediary
symptoms



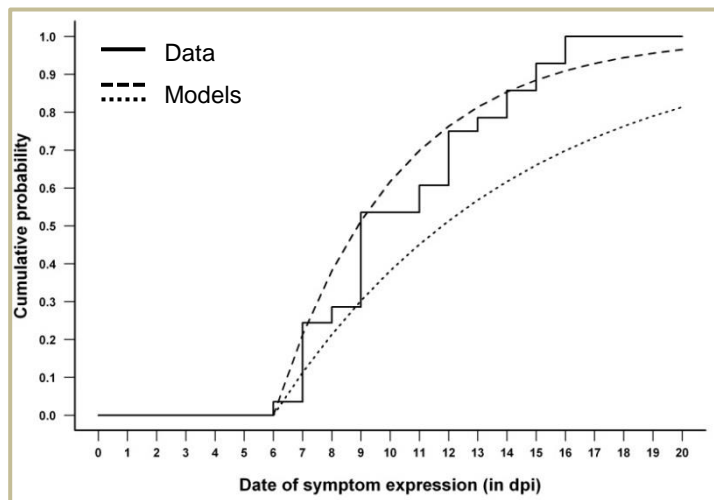
3
Severe
symptoms



4
Highly severe
symptoms

Validation of the protocol by simulation

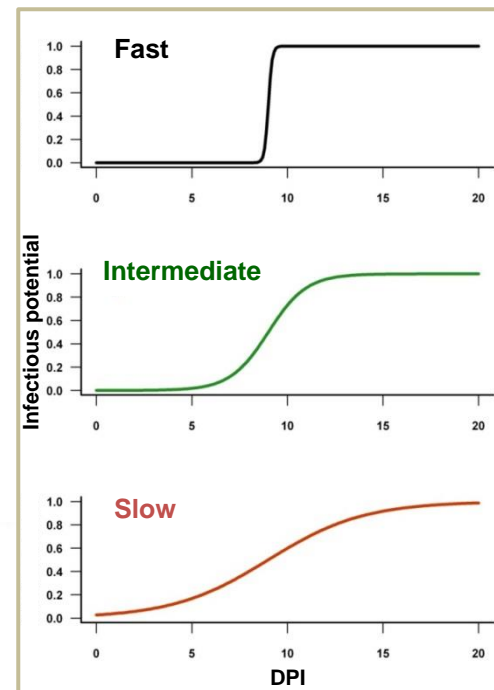
2 symptom expression dynamics to simulate the incubation period



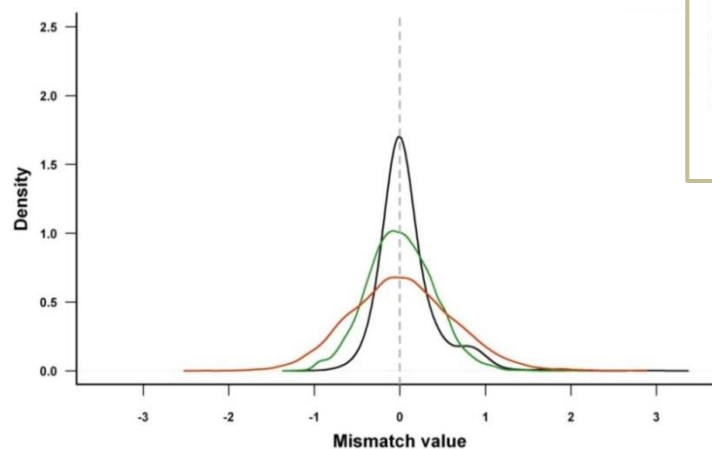
3 scenarios for the mismatch value



3 transition speeds to simulate the latency period



Unbiased and precise estimate of the mismatch



Binomial generalized linear model

$$N_{\text{infected}} \sim B(n_{\text{tested}}, \tau)$$

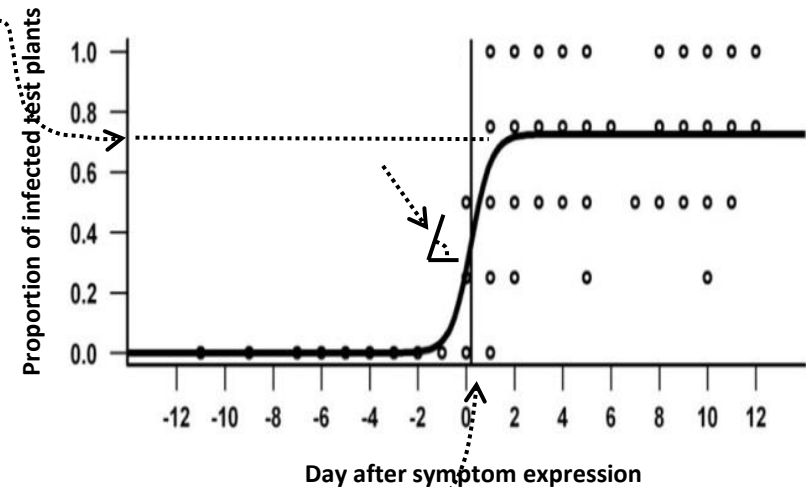
Transmission rate

Maximum transmission rate
(plateau)

$$\tau = \frac{\tau_{\max}}{1 + e^{-\frac{4 \cdot s_m}{\tau_{\max}}(\Delta t - m)}}$$

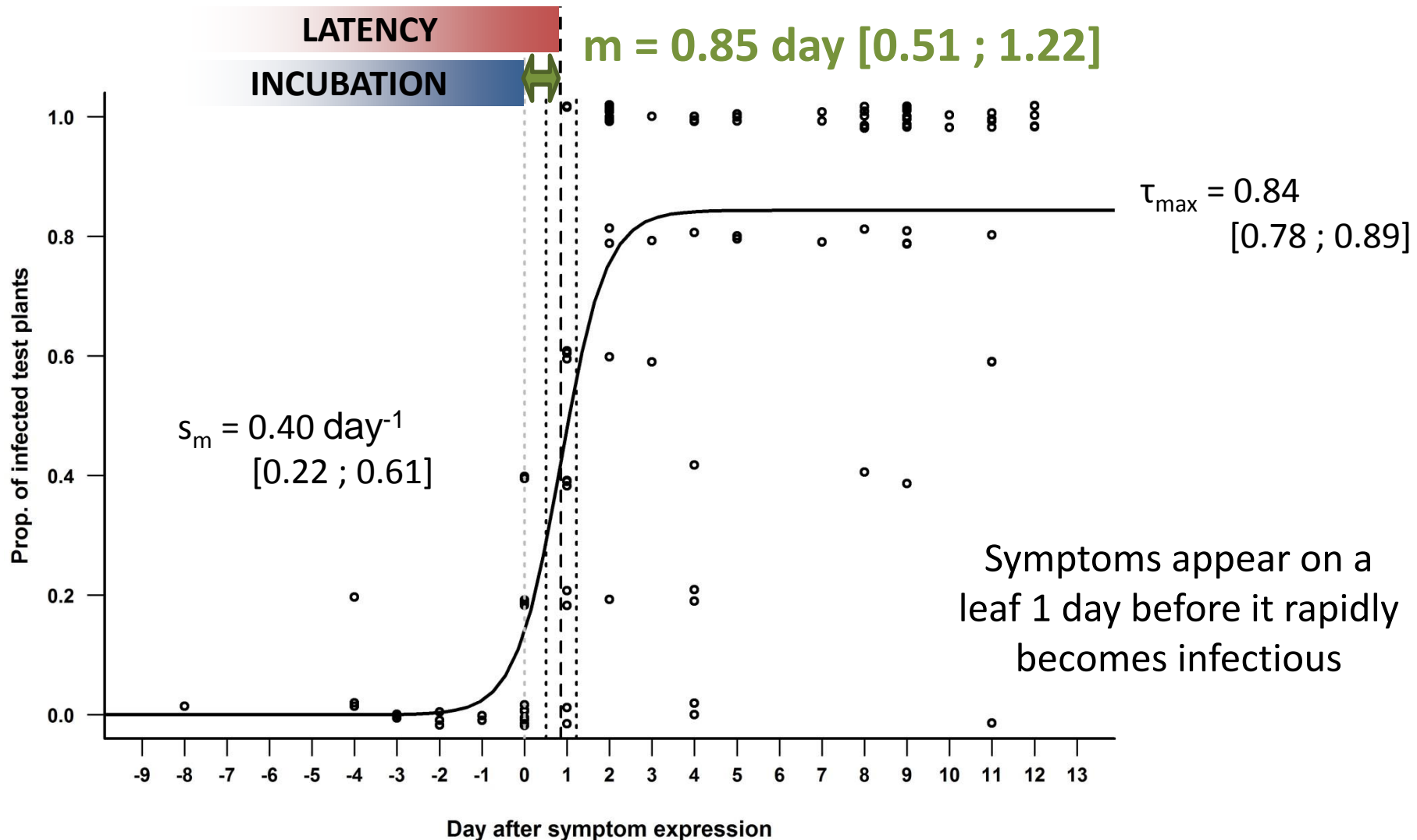
Transition speed
(slope at the inflection point)

Mismatch between incubation and latency
(abscissa of the inflection point)



Parameters estimated by maximum likelihood

Result: a 0.85 day mismatch at the leaf scale



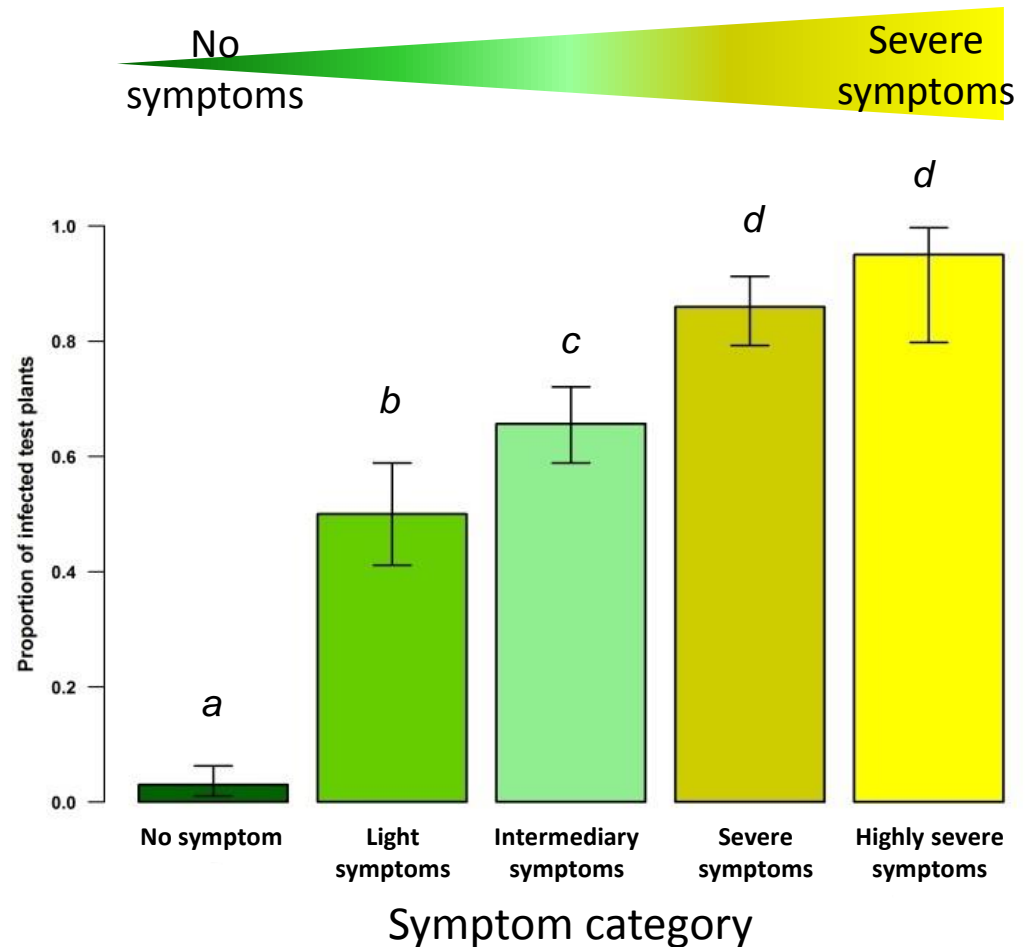
A correlation with symptom severity

Binomial generalized linear model:

The more severe the symptoms, the higher the transmission rate

$$R^2_{\text{McF}} = 0.41$$

Vertical bars: $\text{CI}_{95\%}$



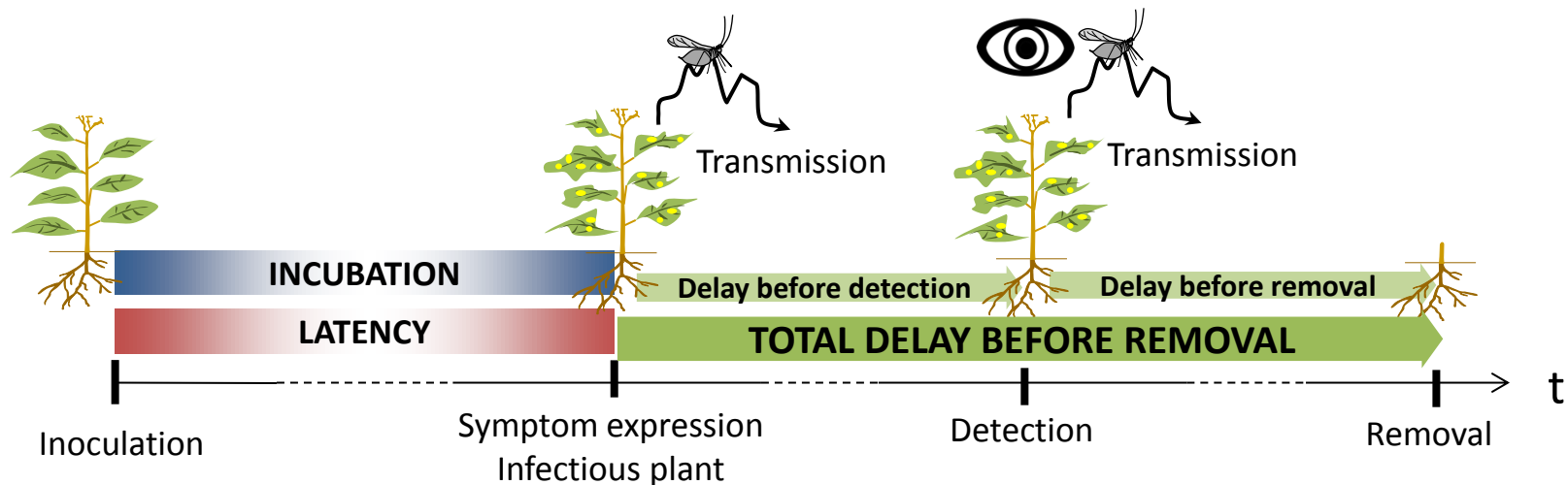
Conclusions

- ❖ A generic experimental approach has been developed to assess precisely the mismatch between incubation and latency
- ❖ Symptom severity and transmission rate are correlated
 - ✓ As suggested for *Cucumber mosaic virus* on *Cucurbita pepo* (Zitter & Gonsalves, 1990)
Cauliflower mosaic virus on *Brassica rapa* (Doumayrou et al., 2012)
- ❖ Under our experimental conditions, latency and incubation of PPV infection of young peach plants are almost synchronized
 - ✓ Symptomatic plants are efficient sources of PPV (Manachini et al., 2004; Damsteegt et al., 2007; Moreno et al., 2009)
 - ✓ *Beet mosaic virus*: latency shorter than incubation of 1 day in *Beta vulgaris* (Dusi & Peters, 1999)

A strategy based on visual detection of plants infected by PPV could be efficient if symptoms are detected without delay?

Future works

- ❖ Modeling the epidemiological impact of the delay between symptom expression and tree removal



- ❖ Development of an early diagnosis procedures
→ Detect an infection before symptom expression

Thank you for your attention

Acknowledgments:

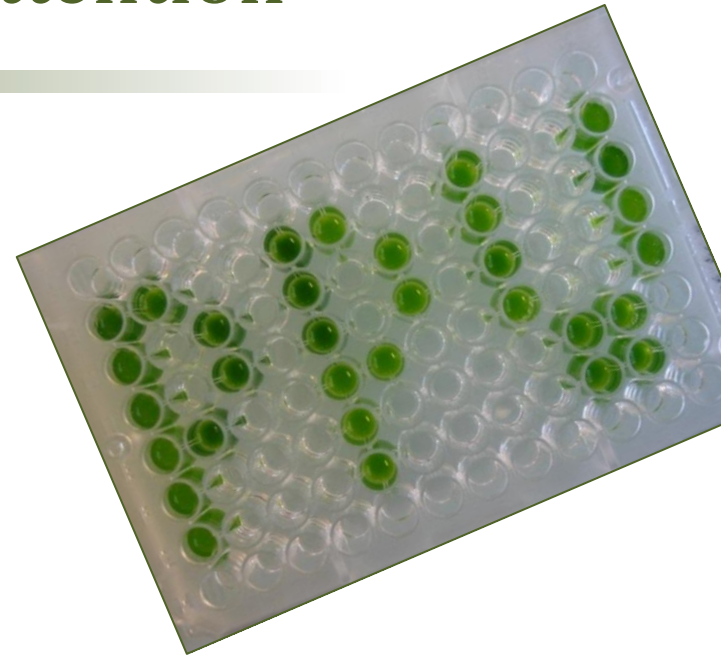
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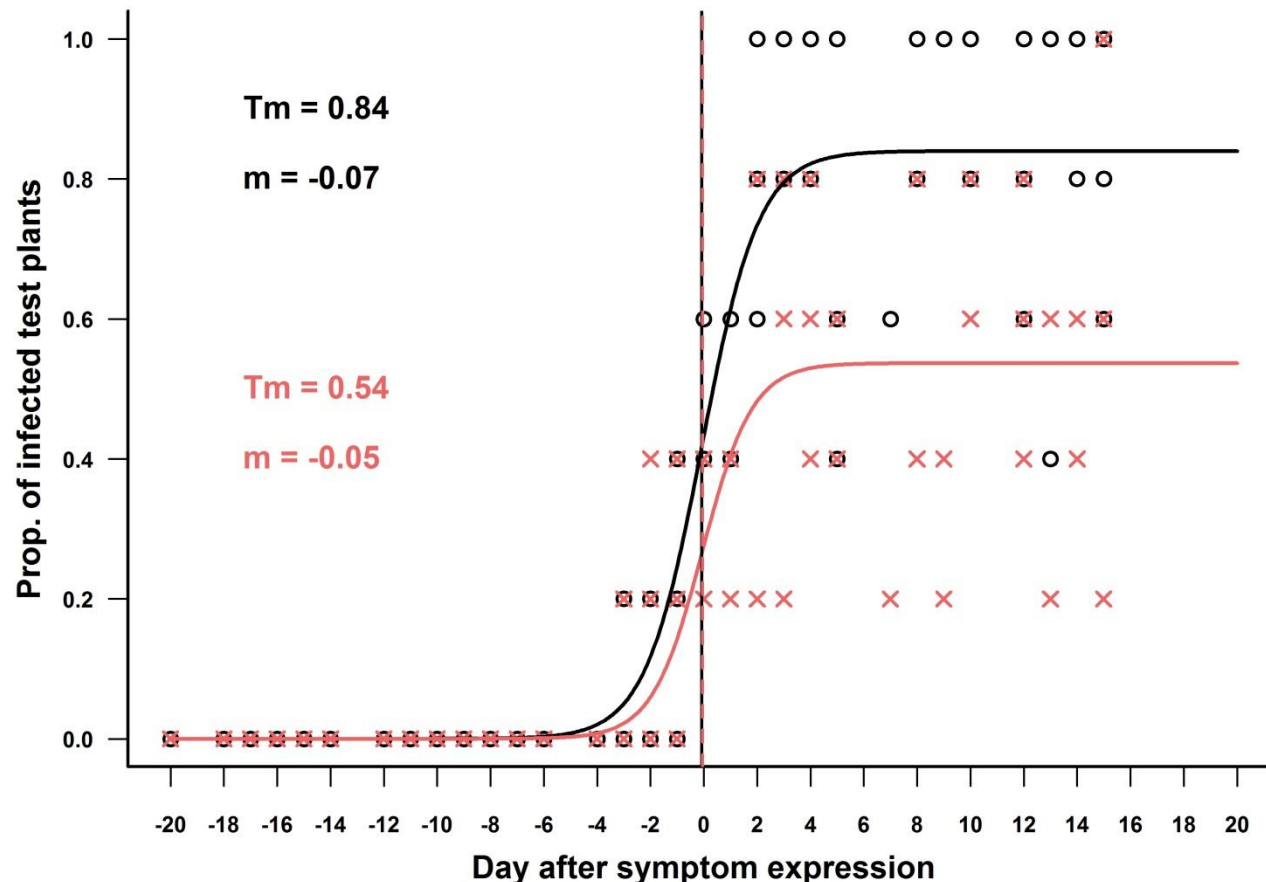
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Bonus: impact of the maximum transmission rate

(simulated experiment)



The abscissa of the inflection point is a robust estimator of the mismatch between incubation and latency

Bonus: generalization to older trees

