## **EVOLUTION OF DECEIT AS A SECOND-LINE DEFENSE:**

 $H_{t+1} = H_t e^{r-aPt}$  $P_{t+1} = H_t (1-e^{-aPt})$ 





# A QUESTION POSED:

# Could a mutant fake marker invade of population of "honest" caterpillars?



# **REFINE QUESTION**

Could a mutant fake marker invade of population of "honest" caterpillars that employ hiding as a first line of defense?



## **SCENARIO**



# **THE CHROMOSOME:** 2 loci, 4 alleles



#### **False Mark**

- High
- Medium
- Low
- No

- Strong
- Medium
- Weak
- None

## THE CHROMOSOME

A binary string that represents the phenotype that expresses a specific strategy:

- e.g. 1100
- 1 1 translates into good hider
- 0 0 translates into no mark



## AN EXAMPLE

Strategy 0010 has a value of 2
i.e. 0\*1 + 1\*2 + 0\*4 + 0\*8

Strategy 1001 has a value of 9 i.e.
 1\*1+0\*2+0\*4+1\*8



## SIMULATION

- Deterministic
- Some proportion of hosts are found based upon relative hiding value
- Some proportion of found hosts are attacked based upon mark strength and wasp rules



#### **UNLIKE PREY-**

#### Hosts don't disappear after attack but they may change identity



# FITNESS IS DETERMINED BY:

- Escape from parasitoids
- Cost of defense
   Hiding≥ false-mark



# **THREE SCENARIOS:**

Parasitoids are hard wired (frequency independent)

Parasitoids are flexible

Parasitoid's flexibility evolves





## **STARTING POPULATION**





## **FREQUENCY INDEPENDENT**





## PARASITOID ACCEPTANCE PLASTICITY

- Within generation as number of "parasitized" hosts increases
- Among generations as fake marker phenotypes change in frequency





## **PLASTIC PARASITES**



![](_page_14_Picture_2.jpeg)

## THE NEW GA

- Actually two GA's, one for the host population and one for the parasitoid population
- The parasitoid chromosome has one gene for sensitivity to mark strength and another for sensitivity to mark frequency in the environment

![](_page_15_Picture_3.jpeg)

#### **PARASITES EVOLVE**

$$FIT(p) = \frac{a\left(Hf_{h} + \sum_{n=1}^{N} Ff_{h}o_{p,n} + Df_{d}o_{p,d}\right)}{1 + a\left(\frac{H}{t_{o}} + \frac{\sum_{n=1}^{N} Fo_{p,n}}{t_{o}} + \frac{\sum_{n=1}^{N} F(1 - o_{p,n})}{t_{r}} + \frac{\sum_{n=1}^{N} Do_{p,d}}{t_{o}} + \frac{\sum_{n=1}^{N} D(1 - o_{p,d})}{t_{r}}\right)}$$

![](_page_16_Picture_2.jpeg)

# **PARASITOID'S PAYOFF**

#### ACCEPT

- FIT(a | p) =  $fp t_o$
- FIT( $a \mid h$ ) =  $fh t_o$

#### REJECT

• FIT =  $-t_r$ 

![](_page_17_Picture_6.jpeg)

#### PARASITES EVOLVE

![](_page_18_Figure_1.jpeg)

![](_page_18_Picture_2.jpeg)

## **AN INSIGHT**

False marking as a second line of defense only spreads through the prey population under limited circumstances

(our test is conservative!)

![](_page_19_Picture_3.jpeg)

#### HOST DYNAMICS

$$\gamma_n(t+1) = \gamma_n(t)(1-\mu)$$
  
where  $\mu = \gamma_n \alpha_n \frac{1}{\sum_{n=1}^N \gamma_n \alpha_T} \rho\left(\frac{m}{T} \beta_n, \sigma\right) \theta$ 

where  $\alpha_n$  = hide genotype n,

- $\frac{m}{T}$  = fraction of marked hosts in population T,
- $\beta_n$  = false mark genotype n,
- $\rho$  = parasitoid response
- $\sigma$  = parasitoid response shape pararameter
- $\theta$  = proportion of hosts encountered per epoch

![](_page_20_Picture_8.jpeg)