Modeling the Evolution of Mimicry

Islam, Grogono

Outline

Batesian Mimicr Mullerian Mimic Evolutionary Dynamics Mimicry Ring

The Model
Past Work
FormAl

Prey Predator

Predator

Experiments

Conclusion

Modeling the Evolution of Mimicry

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The Inspiration

Batesian Mimicry Mullerian Mimicry Evolutionary Dynamics Mimicry Ring

The Model

Past Work FormAl

Prey

Predator

Results

Experiments Analysis

Conclusion

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

He Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

The Model

Prey

Predator

esults

xperiments nalysis

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

The Inspiration

Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

The Model

Predator

Predator

Results

Experiments

onclusion



The Inspiration: Mimicry

The Inspiration: Mimicry

History

Henry W. Bates first published in 1862.

- ► **Content:** Similarity and dissimilarity between Heliconiinae and Ithomiinae butterflies.
- Bates collected 94 species of butterfly.
- Grouped according to similar appearance.
- ▶ Discovery: Appearance: similar, Morphological feature: different species.
- ▶ 67 of 94: Ithomiinae.
- ▶ 27 of 94: Heliconiinae.

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Outline

The Inspiration

Batesian Mimicr Mullerian Mimic Evolutionary Dynamics Mimicry Ring

ie Model

FormAL Prey

Predato

Results

Analysis

The Inspiration: Mimicry

Batesian Mimicry

- Heliconiids are,
 - conspicuously colored
 - extremely abundant
 - slow in mobility.
- Predators, insectivorous birds do not prey on Heliconiids.
 - Reason: Inediblity and unpalatability.
- Heliconiids are easily recalled by predators.
 - ▶ Reason: Conspicuous coloration.
 - Color acts as warning.
- Ithomiinae and Pieridae are,
 - edible
 - palatable
 - pretend like Heliconiids, enjoy protection.

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Outline

Batesian Mimicry

Mullerian Mimicry Evolutionary

Evolutionary Dynamics Mimicry Ring

The Model

Prey

Predator

Results

Experiment



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Batesian Mimicry

According to Wilcker:

Actor is a mime.

False representation of warning pattern: Mimicry.

▶ Bates: First to point out, so Batesian Mimicry.

► **Model:** Animal avoided by predator for unpalatable behavior.

Mimic: Imitating animal.

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Outline

Batesian Mimicry

Mullerian Mimicry Evolutionary Dynamics

The Model

FormAL Prey Predator

Reculte

Experiment



Batesian Mimicry

Plate from Bates (1862)



Figure: Plate from Bates (1862) illustrating Batesian mimicry between Dismorphia species (top row, third row) and various Ithomiini (Nymphalidae) (second row, bottom row).

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Outline

Batesian Mimicry

Mullerian Mimicr Evolutionary Dynamics Mimicry Ring

The Model

Prey Predator

Predator

esuits Experiment:

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Mullerian Mimicry

Two inedible unrelated butterfly species have similar appearance.

Bates: Unable to explain.

Explanation: from Fitz Muller in 1878.

Muller's research was also in Brazil.

Explanation:

- Predator's limited memory.
- ▶ Inedible species loose number.
- Save loss and survival of species:
 - ▶ inedible, different family
 - evolve to have similar appearance.
- Phenomenon: Mullerian mimicry, named after Fritz Muller.

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Outline

The Inspiration
Batesian Mimicr

Mullerian Mimicry Evolutionary Dynamics

Dynamics Mimicry Ring

Past Work

FormAL Prey Predator

esults

xperiments Inalysis



Mullerian Mimicry

Viceroy and Monarch



Figure: A very well-known example of mimicry. Viceroy (top). Unpalatable Monarch (bottom). Image source: Wikipedia

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Outlin

The Inspiration
Batesian Mimicry
Mullerian Mimicry

Evolutionary Dynamics Mimicry Ring

The Mode

FormAL Prey Predator

Predator

Experiment: Analysis



Evolutionary Dynamics

Punctuated Equilibrium

Most sexually reproducing species remain:

- extended state of stasis.
- Little evolutionary change in most geological history.

Cladogenesis:

- Process of speciation.
- One split into two distinct species.
- ► In geological time:
 - Discontinuous change
 - Not gradual transformation

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Outline

The Inspiration

Batesian Mimicry

Mullerian Mimicry

Evolutionary Dynamics

ne Model

Past Work FormAL

Prey Predator

lesults

Experiment: Analysis



Gradually becoming new species.

Phyletic gradualism holds,

- Species population changes gradually.
- No clear demarcation between ancestral and descendant species.
- Gradually changing lineage is divided arbitrarily.
- Evolution is
 - smooth
 - steady
 - incremental
 - but not necessary constant and slow rate in geological time scale.

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Islam, Grogono

Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry

Evolutionary Dynamics

Mimicry Rin

Past Work

Prey Predator

Results Experiment

Analysis Analysis



Evolutionary Dynamics

Punctuated Equilibrium vs. Phyletic Gradualism

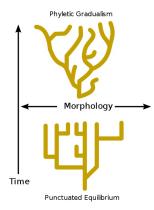


Figure: Punctuated equilibrium (bottom), phyletic gradualism (top). Image source: Wikipedia

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Islam, Grogono

Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry

Evolutionary Dynamics Mimicry Ring

he Model

FormAL Prey

Predator

esuits Experimen

. . .



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Evolutionary Dynamics

- Turner: Synthetic theory.
- Originated from Poulton and Nicholson.
- Mimicry arises in two steps:
 - 1. A comparatively large mutation achieves a good approximate resemblance.
 - 2. A gradual evolutionary change refines the resemblance to a higher degree of perfection.
- Theory also applied to Mullerian Mimicry.

Mimicry Ring

- Examine the local butterfly fauna in any area of the world
 - all the aposomatic species
 - limited number of different patterns
 - normally far smaller than the number of species.
- Mullerian mimicry ring:
 - Each cluster of species
 - all sharing a common pattern
- All the rain forest in South and Central America.
 - most of the long winged butterflies (ithomiids, danaids and heliconids)
 - belong to one of only five different rings.

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Outline

Batesian Mimicry Mullerian Mimicry Evolutionary Dynamics

The Model

FormAL Prey

Predator

Experiments

onclusion

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The Model: Evolution of Mimicry

The Model

Evolution of Mimicry

▶ **Objective:** Build an *agent based* Artificial Life model for simulating the evolution of mimicry.

- ► Two species of agents:
 - 1. Prey
 - Model
 - Mimic
 - 2. Predator
- ▶ Prey pattern representation: Cellular Automata.
- Predator pattern recognition: Hopfield Network.
- ▶ Environment: Visual representation, 3D, toroidal.

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Outline

The Inspiration

Batesian Mimicry

Mullerian Mimicr

Evolutionary

Dynamics

Mimicry Ring

The Model

FormAL Prey Predator

Predator

lesults Experiment

xperiments inalysis



The Model

Past Work

► Turner (1996) and Huheey (1988):

- Focus: Selective pressure on prey, by learning ability of predator.
- Predator: Simple Monte Carlo learning or mathematical approach.
- ► Sherratt (2002):
 - Co-evolving predator and prey population.
 - Predators:
 - deterministic
 - cannot learn from experience.
 - attack policy is fixed, either attack or avoid.

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

The Model

FormAL Prey

Experiment



2. If the Mullerian mimics do not converge into one large ring, then the presence of Batesian mimics could entice them to do so, by influencing the rings to converge.

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicr
Evolutionary
Dynamics
Mimicry Ring

The Model

Past Work

Prey Predator

> Results Experiments



Franks and Noble

Prey and Predator

► Prey:

- Appearance and palatability.
- ▶ Palatability is fixed, using different level from 0 to 1, where 0.5 is neutrally palatable.
- Two genes representing external appearance, using values between 1 to 200.
- Euclidean distance between values represents similarity.

Predator:

- Monte Carlo reinforcement learning system.
- Decision for consumption of prey:
 - ► Use: set of probability formula.
 - ► Dependence on phenotype.

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

ne Model

Past Work

Prey Predator

esults

Experiment



Franks and Noble

Result

- Hypothesis 1 of a single large ring did was not established.
- Hypothesis 2 is established.
- Observation:
 - Batesian mimics provide pressure on mutants.
 - Batesian pressure on mimicry rings push one into the range of another.

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicr
Evolutionary
Dynamics
Mimicry Ring

The Model

Past Work

Prey

Predator

esults

Experiments



FormAl Framework

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FormAl

▶ Ideas from Peter Grogono's Formal Artificial Life (FormAL) project.

Goal:

- Study the emergence of complexity.
- No variable unless genetically controlled or influenced. Principal not followed for Hopfield Network.

Agents:

- Simulated organism.
- Reproduce itself using genetic information.
- Capable of modifying structures of genome between generations.
- Interaction with other agents.
- Survive and reproduce in a challenging environment.

The Model

FormAL Prey

Predator

Results Experimen

Allalysis

Conclusion

Environment is 3D, complete freedom of movement defined from genetic representation.

Space:

- ▶ 3D Lattice of discreet points.
- $(x, y, z) \in \Sigma^3$, where Σ be the set $\{0, 1, ..., S 1\}$.
- S is a universal constant, small positive number. Value is 20, WorldSize parameter.

► Time:

- Integer value, increases in discreet steps.
- Each agent updates itself in each time step.

► Cell:

- ▶ Entire environment is divided into multiple cells.
- ► A 3D cubical section of the hyperspace.
- ► *ISize* parameter kept at 6. Total *ISize*³ = 216 cells.

FormAL Framework

Environment - Visual representation - Front

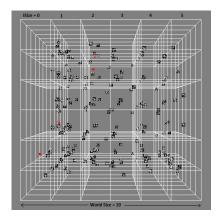


Figure: Three dimensional representation of the environment divided in cells. Presence of different species of agents inside.

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Outline

Batesian Mimico Mullerian Mimico Evolutionary Dynamics Mimicry Ring

The Model

FormAL

Predator

Results

Experiment Analysis



FormAL Framework

Agent Mobility

- Position is calculated in each step in time.
- Vector components: position, force, acceleration and velocity.
- force is calculated from the mobility genes.
- ▶ Newton's law of motion, used to calculate *acceleration* when *force* is not zero.
- acceleration is integrated to obtain velocity and new position.

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

The Model

FormAL Prey

Predator Results

xperiments Inalysis



The Prey

Mimics and Models

- Agent in the FormAL environment.
- Genetic representation of pattern with Cellular Automata.
- Creates diversity of species.
- Pattern evolution is in the process of punctuated equilibrium.
- Mobility and reproduction capability controlled genetically.

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

Past Work

Prey Predator

esults

xperiments

Pattern Representation

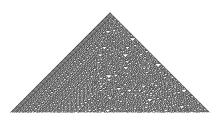


Figure: Cellular Automata Rule 30. Image source: Wikipedia

Current Pattern	111	110	101	100	011	010	001	000
New state of	0	0	0	1	1	1	1	0
center cell								

Table: Cellular Automata rule

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Outline

Fhe Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

ast Work

Prey Predator

Paculte

Experiment:



Batesian Mimicry Mullerian Mimicry Evolutionary Dynamics Mimicry Ring

he Model

Prey

Predator

Results Experiment

Allalysis

- ▶ Pattern genome is 8 bit binary. Decimal range 0 to 255.
- ▶ 256 unique CA pattern.
- Linear representation of pattern stored in Hopfield Network.
- Pattern similarity: Hamming distance between linear representation.
- Single species: group of prey with a specific pattern.
- ▶ Inter species reproduction: restricted to control diversity of patterns.
- "Pattern Mutation Rate": control diversity of new species.

Prey Pattern

Genotype vs. Phenotype

- Genetic bit difference of one.
- Vastly different phenotype.

CA Rule	$60 \equiv 00111100$	$61 \equiv 00111101$	$62 \equiv 00111110$
Pattern	E-2 E-2 E-2 E-2 E-2 E-2 E-2 E-2 E-2 E-2		

Table: Difference in prey pattern genotype and phenotype

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Outline

Fhe Inspiration
Batesian Mimicry
Mullerian Mimicr
Evolutionary
Dynamics
Mimicry Ring

Past Work

Prey Predator

Predator

sults operiments

Genome

▶ 17 bit prey Genome

Pattern(8)	Palatability(2)	Mobility(6)	Reproduction(1)	
10101101	01	110001	1	

Table: Distribution and purpose of each gene of the 17 bit prey genome.

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Outline

Fhe Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

he Model Past Work

Prey

Predator

Results Experiments

Punctuated Equilibrium

Punctuated Equilibrium:

- inclined to cladogenesis instead of gradualism.
- Turner's emphasis on punctuated equilibrium to explain evolution of mimicry.
- CA pattern evolution: single mutation in the pattern genome.
- ► Change of pattern:
 - not gradual
 - arbitrary discontinuous

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

Past Work FormAL Prey

Predator

esults Experiment:



Mobility

Mobility gene: 6 bits.

Use: calculate the magnitude of force for mobility.

Direction:

Towards collection of maximum prey species in a cell.

Away from concentration of predators.

Move towards the selected cell with the magnitude of force calculated. Modeling the Evolution of Mimicry

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Outline

The Inspiration

Batesian Mimicry

Mullerian Mimicry

Evolutionary

Dynamics

Mimicry Ring

he Model ast Work

Prey Predator

Predator

esuits Experiment



Reproduction

- Begins at "Reproductive Age"
- Capability is decided on 17th bit gene.
- Mate selection is random, but within same cell.
- ► Mate needs to be
 - similar species
 - genetically capable to reproduce
 - reached maturity
- Reproduction process:
 - Single point crossover.
 - ► Two point mutation: different rates
 - Pattern Mutation Rate
 - ► Genome Mutation Rate

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

he Model Past Work

Prey Predator

Results

operiments



The Predator

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

l he Model Past Work

Prey

Predator

esults

Experiments Analysis

Conclusion

Agent in the FormAL environment.

- Provide selection pressure for the evolution of mimicry.
- Equipped with Hopfield Network Memory.
- Mobility and reproduction capability controlled genetically.
- ▶ Unable to represent pattern recognition capability with genome.
- New predators are born with zero memory, as memory is not inherited.

The Predator

Learning

- Predator's interaction objective with prey is consumption.
- Consumption is based on palatability.
- If unable to consume prey is thrown back into environment.
- Store prey pattern into memory with the associated palatability.
- New pattern learned with Hebbian Learning.

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

FormAL

Predator

Results Experiments

Analysis



The Predator

Design of Memory

Input to Memory:

- Each prey has an evolving CA represented by a binary Genome.
- ▶ 2D pattern is serialized to a 1D binary array.
- Binary representation converted to bipolar representation.
- Pattern Recognition with Hopfield Network:
 - Learning: Apply Hebbian learning to calculate weights.
 - Initialization: Input to network initialized with input vector.
 - Iterate Until Convergence: Asynchronous update of each neuron. Input: previous state.
 - ▶ Output: Finally a pattern is set as output when the network reaches convergence.

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Outline

he Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

Past Work FormAL

Predator

Results Experiments

Analysis



Predator

- Agent reaches 'Minimum Attack Age' it starts hunting.
- Select a random prey within vicinity (same cell).
- Involves recognition of prey pattern.
- Pattern memorization and recognition process: computationally expensive.
- Two parameters to limit
 - Hopfield Minimum Memory Size (value 2 to 6)
 - Hopfield Maximum Memory Size (value 10)
- New predator attacks without caution.
- Attacks everyone and in the process store pattern and palatability.
- ▶ When memory reaches 'Hopfield Minimum Memory Size': intelligent selection.

The Predator

Genome

Mobility(4)	Reproduction(1)		
1101	1		

Table: Distribution and purpose of each gene of the 5 bit predator genome.

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Outline

The Inspiration

Batesian Mimicry

Mullerian Mimicry

Evolutionary

Dynamics

Mimicry Ring

e Model ast Work ormAL

Prey Predator

Predator

esults

operiments

The Predator

Mobility

- ▶ Magnitude of force of movement: 4 bits of genome.
- Direction: number of prey species present in a neighboring cell.
- Move towards the cell with least number of predator and most number of prey.
- Result: Predators are distributed all over the environment.
- Increases agent's predatory behavior.

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Outline

The Inspiration

Batesian Mimicry

Mullerian Mimicry

Evolutionary

Dynamics

Mimicry Ring

FormAL Prey

Predator

esults experiments



The Predator

Reproduction process

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Modeling the

Evolution of

Outline

- ► Reproduction age: begins.
- ▶ Reproduction age interval: which is difference in time between two reproduction activity.
- Capability: 5th gene.
- Selection: another predator from same cell which satisfy above conditions.
- Perform genome crossover and mutation: create new predator.
- New predator initialized with zero memory.

The Inspiration
Batesian Mimicry
Mullerian Mimicr
Evolutionary
Dynamics

Past Work FormAL

Prey Predator

Danulka

experiments



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Islam, Grogono

Outline

The Inspiration

Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

The Model

FormAL Prey

Predator

Results

Experiments Analysis

Conclusion

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The Results

Results

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Results

Objective: Evaluate evolution of mimicry.

- Evaluation process:
 - Calculate number of mimicry rings.
 - Calculate size of mimicry rings:
 - Population of palatable species.
 - Population of unpalatable species.
- Report parameters:

Parameter	Value			
Mimicry Ring hamming distance	10 % of the Pattern Size			
Number of Rings to report	8			

Table: Parameters to mimicry ring report.

Initial Configuration

	1				
	Prey configuration			Predator configuration	
Population	Rule110 (Palatable)	Carr	108	10	
	Rule30 (Unpalatable)		108		
Reproduction	Age Limit	100		500	
	Interval	1000		1200	
Mutation Rate	Pattern	0.05		0.3	
	Genome	0.5			
Demise Age	2000			2500	
Minimum Attack Age			500		
Memory Configuration				Minimum	2
				Maximum	10

Table: Agent configuration of 2 prey species

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

he Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

The Model

FormAL Prey Predator

Results

Experiments Analysis

Population vs. Time (10k)

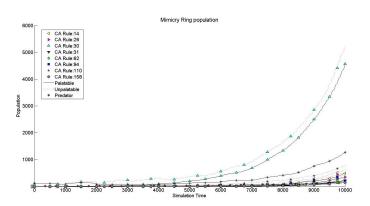


Figure: Population distribution of mimicry rings, initialized with 2 prey species, 10k iterations

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Birch

The Model

FormAL Prey Predator

Result

Experiments Analysis

Population vs. Time (5k)

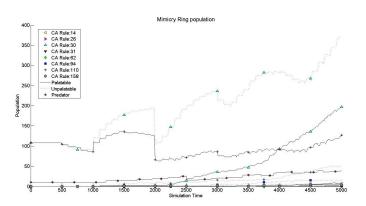


Figure: Population distribution of mimicry rings (2 prey species, 5k iterations)

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics

Past Work FormAL Prey

Predator

Experiments

Analysis

Number of Mimicry Rings

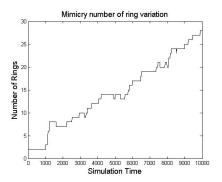


Figure: Number of mimicry rings, initialized with 2 prey species.

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Bias

he Mode Past Work FormAL

Prey Predator

Results

Experiments

Generated Prey Species

Population vs. Time (10k)

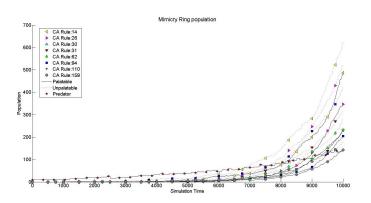


Figure: Population distribution of generated mimicry rings (2 prey species)

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Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics

The Model Past Work

Prey Predator

Result

Experiments Analysis

Four Prey Species

Increased Population

	Prey configuration			Predator configuration	
Population	Rule110 (Palatable)		150 ↑	20 ↑	
	Rule30 (Palatable)		150 ↑		
	Rule55 (Unpalatable)		150 ↑		
	Rule190 (Unpalatable)		150 ↑		
B 1 2	Age Limit	100		500	
Reproduction	Interval	1000		2500 ↑	
Mutation Rate	Pattern	0.05		0.3	
Mutation Rate	Genome	0.5			
Demise Age	2000			7000 ↑	
Minimum Attack Age			500)
Memory Configuration				Minimum	4
				Maximum	10

Table: Agent configuration of 4 prey species with increased population

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Outline

Fhe Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics

The Model

ormAL Prey Predator

Results

Experiments Analysis



Four Prey Species

Increased Population Population vs. Time (10k)

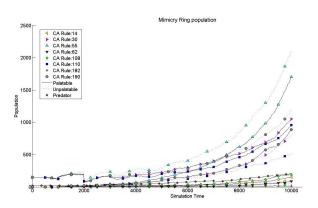


Figure: Population distribution of mimicry rings (4 prey species, increased population)

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Outline

Fhe Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

The Model Past Work

Prey Predator

Results

Experiments Analysis



Only Unpalatable Species

Initial Configuration

	Prey configuration			Predator configuration	
	Rule110 (Unpalatable)		150	20 ↓	
Population	Rule30 (Unpalatable)		150		
	Rule55 (Unpalatable)		150		
	Rule190 (Unpalatable)		150		
Reproduction	Age Limit	100		500	
Reproduction	Interval	1000		2000	
Mutation Rate	Pattern	0.05		0.3	
	Genome	0.5			
Demise Age	2000		5000 ↓		
Minimum Attack Age			500		
Manage Confirmation				Minimum	4 ↓
Memory Configuration				Maximum	10

Table: Agent configuration of 4 prey species all unpalatable.

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Outline

The Inspiration

Batesian Mimicry

Mullerian Mimicry

Evolutionary

Dynamics

Mimicry Ring

Past Work FormAL

Prey Predator

Result

Experiments Analysis

Only Unpalatable Species

Population vs. Time (10k)

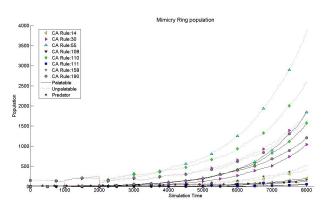


Figure: Population distribution of mimicry rings(4 prey species all unpalatable)

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

he Mode Past Work

Prey Predator

Results

Experiments Analysis

Only Unpalatable Species

Reduced Predator Memory Population vs. Time (10k)

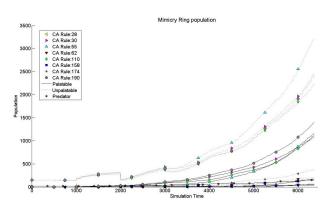


Figure: Population distribution of mimicry rings. 4 prey, all unpalatable but reduced predator memory

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

Fhe Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

Past Work FormAL Prey

Prey Predator

Experiments

Experiments Analysis



Only Palatable Species

Initial Configuration

	Prey configuration			Predator configuration	
Population	Rule110 (Palatable)		150		
	Rule30 (Palatable)		150	20	ı
	Rule55 (Palatable)		150		
	Rule190 (Palatable)		150		
D I	Age Limit	100		500	
Reproduction	Interval	1000		2000	
Mutation Rate	Pattern	0.05		0.3	
	Genome	0.5			
Demise Age	2000			5000	
Minimum Attack Age				500	
M				Minimum	4
Memory Configuration				Maximum	10

Table: Agent configuration of 4 prey species all palatable.

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Islam, Grogono

Experiments

Only Palatable Species

Population vs. Time (7k)

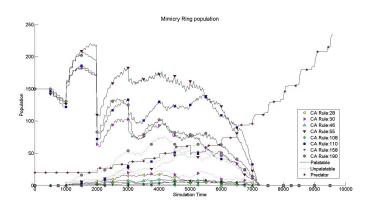


Figure: Population distribution of mimicry rings (4 prey species all palatable)

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

Past Wor FormAL Prey Predator

Result

Experiments Analysis

Only Palatable Species

Number of Mimicry Rings

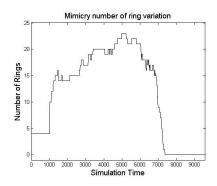


Figure: Number of mimicry rings (4 prey species all palatable)

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

The Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

Past Work FormAL Prey

Predator

Experiments

Analysis

Analysis

Batesian Mimicry

- Batesian Mimicry has taken effect, for all possible initial conditions.
 - Every ring of unpalatable species there is a palatable ring.
- Start with palatable population, prey reaches extinction.Reason: No models to mimic for palatable species.
- Conclusion: This model can simulate evolution of Batesian Mimicry.

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

Fhe Inspiration
Batesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

Past Work FormAL Prey Predator

Results
Experiments
Analysis



Analysis

Mullerian Mimicry

"Mullerian mimics converge into one large ring."

- Initialize simulation with 4 unpalatable species. No palatable ones.
- After 10k iteration all unpalatable species survive with dominance.
- ▶ **Reason:** Predator minimum memory configuration set to 4.

New experiment:

- Reduce predator memory to 1.
- ▶ **Observation:** Single large ring do not occur.
- Conclusion: Similar to Franks and Noble. Multiple Mullerian mimics do not converge into one large ring.

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

he Inspiration
Satesian Mimicry
Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

he Model Past Work

Prey Predator

Predator

Experiments

Analysis



Analysis

Conclusion

- Successful simulation of evolution of mimicry.
- Accurate simulation of mimicry ring. Diverse new rings and shift in their population.
- ▶ Proof of the theory of Turner: evolution of mimicry with punctuated equilibrium.

Modeling the Evolution of Mimicry

Islam, Grogono

Analysis

Modeling the Evolution of Mimicry

Islam, Grogono

Outline

The Inspiration

Mullerian Mimicry
Evolutionary
Dynamics
Mimicry Ring

The Mode

FormAL Prey

Predator

Results

Experiments Analysis

Conclusion



Conclusion

Modeling the Evolution of Mimicry

Islam, Grogono

Conclusion

Artificial Life:

- Tool for biological inquiry
 - Success: Proof of Turner's punctuated equilibrium
 - Proof of Franks and Noble's converge of one large ring.
- Nature inspired computer science
 - Success: Appropriate emulation of Batesian and Mullerian mimicry.
- Solve problem in computer science
 - ▶ Not successful: unable to find the appropriate problem solving scenario.