Adapting genetic algorithms for artificial evolution of visual patterns under selection from wild predators

Short title: Camouflage evolution in the wild

Emmanuelle S. Briolat[1], George R. A. Hancock[1] & Jolyon Troscianko[1]

[1]: Centre for Ecology and Conservation, Faculty of Environment, Science and Economy, University of Exeter, Penryn, TR10 8ER, UK

Corresponding author:

E. S. Briolat, esb204@exeter.ac.uk

Centre for Ecology and Conservation, University of Exeter, Penryn, TR10 8ER, UK

Abstract

Camouflage is a widespread and well-studied anti-predator strategy, yet identifying which patterns provide optimal protection in any given scenario remains challenging. Besides the virtually limitless combinations of colours and patterns available to prey, selection for camouflage strategies will depend on complex interactions between prey appearance, background properties and predator traits, across repeated encounters between co-evolving predators and prey. Experiments in artificial evolution, pairing psychophysics detection tasks with genetic algorithms, offer a promising way to tackle this complexity, but sophisticated genetic algorithms have so far been restricted to screen-based experiments. Here, we present methods to test the evolution of colour patterns on physical prey items, under selection from wild predators in the field. Our techniques expand on a recently-developed open-access pattern generation and genetic algorithm framework, modified to operate alongside artificial predation experiments. In this system, predators freely interact with prey, and the order of attack determines the survival and reproduction of prey patterns into future generations. We demonstrate the feasibility of these methods with a case study, in which free-flying birds feed on artificial prey deployed in semi-natural conditions, against backgrounds differing in three-dimensional complexity. Wild predators reliably participated in this experiment, foraging for 11 to 16 generations of artificial prey and encountering a total of 1,296 evolved prey items. Changes in prey pattern across generations indicated improvements in several metrics of similarity to the background, and greater edge disruption, although effect sizes were relatively small. Computer-based replicates of these trials, with human volunteers, highlighted the importance of starting population parameters

23   for subsequent evolution, a key consideration when applying these methods. Ultimately,

24   these methods provide pathways for integrating complex genetic algorithms into more

25   naturalistic predation trials. Customisable open-access tools should facilitate application of

26   these tools to investigate a wide range of visual pattern types in more ecologically-relevant

27   contexts.

Keywords

30   camouflage; evolution; genetic algorithms; predation; selection; virtual ecology

Introduction

Camouflage is a common anti-predator defence strategy, with a long history as a case study

for evolution [1]. A vast body of work has investigated how visual patterns help conceal prey

from predators, identifying a range of strategies from disruptive coloration and

countershading to masquerade and iridescence [1,2]. Understanding which patterns

produce the most effective protection is an important goal for camouflage research,

particularly with respect to practical applications, yet determining which combinations of

traits provide optimal camouflage is not straightforward. The typical approach of comparing

the detectability of a set of phenotypes, varying in specific traits, has been hugely successful

in determining relevant features for crypsis, but is poorly-suited to exploring camouflage

optimisation: the vast phenotypic space available for colours and patterns, along with

interacting effects of viewing conditions, backgrounds and receiver vision, make it

unfeasible to test all possible combinations in this way [3]. Moreover, prey patterns are

shaped by long-term interactions between communities of predators and prey, learning

across repeated encounters and co-evolving over evolutionary time. In particular, search

image formation, whereby predators learn to more efficiently detect common prey types, is

thought to lead to apostatic selection, preventing the evolution of a stable optimum and

favouring polymorphisms in prey patterns [4,5].

Inspired by the processes of natural selection, artificial evolution of visual patterns using

genetic algorithms presents an alternative approach, allowing for exploration of a wide

phenotypic space and accounting for the effects of repeated interactions between

predators and prey. Originally developed as a general solution to optimisation problems

with vast parameter spaces [6], genetic algorithms are an increasingly popular technique in

biological research, ideally suited to exploring visual pattern space. So far, genetic

algorithms have been used to investigate the evolution of several aspects of visual

signalling, primarily camouflage patterns [4,7–12], but also aposematism [13], motion dazzle

[14] and even nectar guides in flowers [15]. In brief, genetic algorithms set out both virtual

genetic structures underpinning phenotypes and rules governing survival and reproduction

into the next generation, based on the responses of an observer; across multiple encounters

with the observer in successive generations, optimal solutions to the task should arise.

When the observer is a foraging predator, prey items that are easiest to detect are selected

against, and so lost from the population, while those that are most difficult to find will

survive and breed, ultimately leading to the evolution of better-camouflaged patterns.

Following similar principles, early experiments in virtual ecology tracked populations of

different-coloured pastry baits under selection from wild birds, with the proportion of

different bait types in each generation determined by the ratio of survivors in the previous

round [16,17]. Building on this simple paradigm, genetic algorithms incorporate processes of

mutation and sexual reproduction, allowing for more complex variation in prey phenotype,

and for novel traits to arise. In addition to the algorithms determining how patterns are

retained, lost or modified between generations, the ways in which different pattern traits

are encoded is also increasingly sophisticated. While no models claim to replicate the actual

genetic architecture of a particular species, advances in our understanding of how animal

patterns are genetically determined and physically produced during development [18] have

inspired systems better able to produce a range of realistic natural patterns [12].

So far, these more complex genetic algorithms have generally been applied only in

computer-based experiments, where they are most easily implemented, but which fail to

capture the impacts of naturalistic viewing conditions and relevant receiver vision.

Techniques such as projecting the search area on larger screens, using natural background

images with occlusion layers, or simulating the visual system of dichromats can increase the

realism of the search task [11], but still fall short of replicating the effects of real predator

behaviour. To our knowledge, genetic algorithms have only been combined with selection

by non-human predators in one scenario: seminal experiments on frequency-dependent

selection in which blue jays (*Cyanocitta cristata*) searched for artificial moths with evolving

patterns on screens [4,7]. Besides restricting prey patterns to greyscale targets, these

experiments required highly-motivated captive predators - undergoing up to eight months

of pre-training then 100 generations' worth of trials [7] - so are difficult to expand to other

species and contexts.

Here, we describe a more generalisable process that combines the benefits of field trials

involving natural predators with state-of-the-art tools for artificial evolution, expanding on a

recently-released genetic algorithm and pattern generation framework, the CamoEvo

toolbox [12]. Prey patterns are generated by CamoEvo, with modifications facilitating

implementation in the field, and deployed on physical prey items, concealing a food reward.

Wild predators then interact with these items in semi-controlled arenas, and the order in

which prey are attacked guides the evolution of subsequent generations of patterns. As a

proof-of-principle, we present a case study in which we tested for effects of background

complexity on the evolution of camouflage patterns, in artificial prey under selection from

wild birds. We also replicated the field trials in computer-based experiments with human

volunteers, to explore how field experiments could be rapidly prototyped and parametrised

in future applications.

Materials and Methods

General principles

We developed a methodology for generating and testing artificial prey in the field using

genetic algorithms, by substantially adapting an existing screen-based implementation, the

CamoEvo toolbox [12], released here as CamoPrint in CamoEvo V2.0 (see supporting code).

The properties of the prey and predation scenarios can be customised, and multiple colour

and pattern metrics can be used to analyse the outcomes of pattern evolution, providing

flexibility to suit a range of research questions (Fig 1). To begin with, an initial population of

prey patterns is created by the CamoPrint software, following experimenter inputs for

starting parameters, such as prey size, shape, population size and the colour space from

which prey colours can be drawn (see [12] and supporting code). This first generation of

patterns is applied to real physical objects and deployed where predators of interest can

interact with them. Suitable prey items can take virtually any form required, so long as they

can conceal a food reward accessible to predators, or otherwise provide a record of

predation attempts. We describe a technique for transferring computer-generated patterns

onto plasticine or 3D-printed shapes (see case study), but appropriate two- and three-

dimensional targets can also be made more simply by printing on paper (Fig 1). The setting,

visual backgrounds and predators recruited will depend on the experimental question; these

methods could be implemented in a wide range of scenarios, from relatively controlled

predation trials with captive animals to more naturalistic experiments with multiple

predators and prey scattered across a large area. As predators interact with prey, the timing

or order of predation is recorded, whether through direct observation, filming, or regular

checks, as in more traditional artificial predation experiments. Each population of prey can

also be split into discrete groups, for example if prey are to be presented in a series of semi-

controlled arenas (see case study). In that case, prey can be ranked according to the order

of attack within each group; the size and number of groupings, and prey arrangement within

arenas, can be customised as needed. Predation outcomes are then fed back into the

genetic algorithm software, which uses this information to determine which patterns

survive and reproduce into the next generation. To facilitate the integration of physical prey

items into the existing CamoEvo workflow, we implemented several modifications to the

pattern generation tools – to create the target shapes and shading used in the case study,

assign prey to groups and positions in test arenas, and more easily input the results from

predation trials – which can be adapted to suit each user's needs (see Supporting

Information (SI) for details, and supporting code). Repeating these processes of pattern

generation, prey preparation, predation trials and data entry allows experimenters to follow

selection by predators across multiple generations of prey.

Fig 1: Examples of possible target types and general methods workflow. Users can

customise a range of experimental parameters, and investigate multiple pattern traits when

analysing results. Prey presented here include paper targets, two-dimensional with simple

or more specific shapes (top), and three-dimensional rolled paper tubes, as well as 3D-

printed shells (middle, as used in case study). Rightmost panels show example prey *in situ*.

Finally, to explore the outcome of selection, experimenters can select a wide range of

colour, luminance and pattern metrics to analyse changes in prey appearance across

generations. CamoEvo can analyse the computer-generated prey patterns using a range of

tools in the widely-used MICA and QCPA toolboxes in ImageJ [19,20], providing a fully open-

access workflow, from experimental design to the assessment of pattern evolution.

Alternatively, the physical prey items themselves can be photographed and processed using

the same open-access tools, so that image analyses can account for the real 3D properties

of prey and backgrounds, and relevant visual systems.

Case study - prey pattern evolution against backgrounds of varying three-dimensional

complexity

As a proof-of-concept, we present an application of these methods, investigating the

evolution of camouflage patterns in response to selection from wild birds. We deployed

161 prey in arenas with specific visual backgrounds, to test the effect of differences in three-

162 dimensional background complexity on the strength and direction of pattern evolution.

163 Theoretical models [21] and experiments with both human and avian predators [22–24]

164 suggest even relatively poorly-camouflaged targets are more difficult to detect against

165 complex backgrounds, relaxing selection for increased similarity to the background [24]. In

166 addition, three-dimensional backgrounds introduce variation in lighting conditions, between

167 shadowed areas and those under direct lighting, which creates noise in both luminance and

168 colour [25,26]. As a result, we expected to see improved camouflage in our targets across

169 generations, but with a weaker signal of directional change in the population of prey

170 displayed against backgrounds with greater three-dimensional complexity.

172 *Field set-up*

173 Two sets of artificial predation experiments were run in different locations in the grounds of

174 the University of Exeter's Penryn campus (United Kingdom; 50° 10' 12'' N, 5° 7' 22.8'' W),

175 from 17th September to 21st October and 10th November to 11th December 2021.

176 Experiments were approved by the University of Exeter's ethics committee (Application ID:

177 eCORN003896 v3.1).

179 Our set-up was designed to encourage natural foraging behaviour by wild birds, while

180 allowing us to control the backgrounds prey were viewed against, and accurately track

181 predation events. Prey were placed in six wooden trays (53x53cm,22cm above the ground),

182 set out in areas where birds likely to interact with the set-up were regularly seen. Each tray

was lined with a polypropylene ground cover membrane, allowing water to filter through,

and a mix of gravel and construction sand, forming the visual background (Fig 2). Many

alternative backgrounds could be used here; this gravel mix was selected as a relatively

natural and heterogeneous option, that was also stable enough to remain in place in poor

weather conditions and could be sculpted to vary the complexity of the scene. To test for an

effect of background complexity, each experimental run monitored the survival of two

populations of prey, presented in trays in which the gravel was either smoothed flat or given

3D structure, with wavy furrows creating peaks and troughs in various orientations

(henceforth, smooth and furrowed treatments; Fig 3).

Fig 2: Experimental set-up for field trials. Tray array for the first run (a), with the main

predators involved, *C. corone* (b) and *E. rubecula* (c), and a side view of a tray and

mechanical timing gate (d). Here, the first two items attacked were associated with the

green and red marbles (leftmost prey).

Fig 3: Schematic of the process for selection and evolution of prey patterns in each

generation. Inset: Diagram representing prey positioning in the trays, with strings

connecting to the mechanical timing gate.

Each population consisted of 24 prey, divided into six groups of four per tray. Every

morning, half of each population was placed out into three trays with smooth or furrowed

gravel respectively, so each generation of both populations was fully exposed to predators

across two consecutive days (Fig 3). The location of the smooth and furrowed backgrounds among the trays was alternated daily, and the prey positions in each tray were pseudo-randomised using a custom script in ImageJ [27], designed to reduce spatial clustering. In CamoPrint, custom additions to the CamoEvo toolbox automated the allocation of prey patterns to each tray and labelled them for easier identification (see SI and supporting code). The artificial prey items themselves consisted of 3D-printed oval shells (18x14 mm) which birds had to flip over to access a concealed food reward (dried mealworm and suet). Prey pattern was manipulated by covering the shells in a thin (~1mm) layer of white tack (UHU, Bühl, Germany), then applying custom patterns printed on waterslide transfer paper (Rolurious) with a laser printer (Colour Laser Jet Enterprise M553, HP, USA). Prior to the experimental trials, wild birds were gradually attracted to the trays and trained to search for the artificial prey, starting with a single tray with visible baits, moving on to food in upturned prey shells, then concealed rewards, and finally increasing the number of trays and prey. During training, prey patterns consisted of checkerboards with colours sampled from the colour space of the experimental patterns. Trials began once visiting birds had consumed four prey in all six trays for at least three consecutive days. In the first run, the trays were visited by carrion crows (*Corvus corone*); in the second, a robin (*Erithacus rubecula*) was the main predator, with occasional visits from blackbirds (*Turdus merula*) and great tits (*Parus major*) (Fig 2; S1 Fig; S1 Video).

During experimental trials, the order in which prey items were attacked in each tray served as our measure of prey detectability. To automatically record this, the prey shells were tethered from the underside to coloured strings running under the trays, connecting them

to a mechanical timing gate. The four prey items in each tray were always assigned string

colours in the same order – red, green, blue and yellow from left to right– with marbles of

matching colours in the timing gate; when an item was attacked, its string was pulled,

releasing a marble of the corresponding colour to create a visible record of attack order

(Figs 2,3). This system was not always sufficiently sensitive to record attacks from smaller

birds, so as a fail-safe, each tray was also filmed continuously with a small spy camera

(Coulax).

Prey were set out each morning (8 - 9am), and trays were checked every two to three hours;

each trial ended once at least three items had been attacked per tray. If no prey were

attacked over the course of the day, the trays were covered up overnight, and the same

prey presented again the following day. Once each generation had been fully exposed to

predation, the order of prey attacks in every tray was entered into the genetic algorithm

software, and a new set of prey patterns was produced for the next generation. See Fig 3 for

a full schematic of the experimental procedure.

Pattern generation and algorithm set-up

Prey patterns were generated using the CamoEvo toolbox's animal pattern generator, with a

modified script for the 3D targets used in this experiment (See SI). Targets were oval, with

colourations occupying a CIELAB space with ranges 0 – 100 in luminance, -50 – 50 in A

(green-red) and -10 – 70 in B (blue-yellow). Every population began with an identical set of

patterns, randomly-generated using these parameters with a uniform distribution of genes

(Generation 0; S2 Fig). Details of the pattern generation and genetic algorithm framework

can be found in [12].

CamoEvo is primarily designed for computer-based experiments, so we implemented

several modifications to facilitate its application to a field experiment with physical prey

items, where prey survival must be manually reported (described in full in SI). These

modifications form the basis of the CamoPrint tools featured in CamoEvo V2.0. Once the

trials for each generation were complete, prey were assigned a fitness ranking from 1 to 4 in

every tray, based on the order in which they were attacked. The genetic algorithm then

selected the last two patterns attacked in each tray (ranks 3 & 4) to survive and reproduce,

creating the next generation of patterns. However, to reduce the likelihood of relatively

well-camouflaged prey being eliminated by chance, we applied CamoEvo's built-in rescue

system [12]: if a prey pattern was ranked 1 or 2 in generation N, but ranked 4 (the highest

possible rank) in generation N-1, the pattern also survived into the next generation (N+1). In

the first run, trials stopped after generation 10, when crows stopped visiting the site; in the

second, trials ended as planned after 15 generations. To account for the final set of predator

decisions, we used CamoPrint to produce an additional generation of patterns (generation

11 or 16 respectively), included in subsequent analyses.

Prey camouflage was assessed with digital photography and image analysis, using well-

established methods implemented in the Multispectral Image Calibration and Analysis

(MICA) and QCPA plugins for ImageJ [19,20]. The following commonly-used metrics of

camouflage efficacy were analysed, accounting for perception by relevant avian visual

systems: chromatic (ΔS) and achromatic differences (ΔL) between mean colours in the

Receptor Noise-Limited space [20,28,29], pattern difference based on granularity analyses

[30], and target edge disruption, measured using the GabRat method [31]. To account for

prey pattern, where component colours may better match the background than the mean

prey colour, we also calculated a frequency-weighted average measure of colour difference

between all colours in each prey pattern and the most similar colours in the background.

Statistical analyses were carried out in R version 3.5.2 [32]. See SI for details.

Screen-based search tasks

The outcomes of the field trials suggested our methods could be improved by refining the

experimental parameters, to adjust task difficulty. To investigate this, we replicated the field

trials with matching screen-based search tasks with human volunteers, allowing us to more

rapidly test the impacts of different starting conditions; compared to a month-long run of

field trials, each replicate of the screen-based task took roughly only half an hour for a

participant to complete. These search tasks were also designed using the CamoEvo toolbox,

and run using its built-in user interface for screen-based psychophysics experiments. All

parameters were selected to mimic the field experiment as closely as possible, with the

exception of starting colour space: target patterns occupied either the same colour space as

prey in the field (henceforth the full colour space), or a narrower colour space,

corresponding to 2 standard deviations around the mean colour of the background images.

In the initial generation 0, targets from the full colour space were thus identical to

generation 0 targets in the field, while those from the narrow colour space featured the

same patterns but with a restricted range of colours (S5 Fig). Each participant viewed a

single population of targets over 16 generations, shown on photographs of the same

smooth or furrowed gravel backgrounds as in the field. Every image contained four targets

to find, and participants were tasked with clicking on them as quickly as possible, within a

maximum of 15 seconds per slide. Survival of targets into the next generation and pattern

evolution were governed by the same rules as in the field, based on the order in which

targets were clicked. Changes in the detectability of targets over successive generations

were quantified using capture time, the time taken for participants to click on each target,

as recorded by the CamoEvo toolbox. Measures of prey phenotype used to assess

camouflage efficacy, and statistical analyses, were also selected to match those used for

prey in the field trials as closely as possible (see SI for details of the experimental design and

analyses). In addition, we used CamoEvo to produce 12 replicate populations, with identical

starting conditions to the experimental populations, but in which prey patterns were

selected at random to survive and reproduce into the next generations. These provide

control trends, against which experimental populations can be compared to demonstrate

the effect of selection by human volunteers, as opposed to neutral drift.

A total of 12 volunteers completed these screen-based tasks, between 23rd March and 13th

April 2022 (three per combination of background type [smooth or furrowed] and colour

space [full and narrow]). Experiments took place in a dark room, with images displayed on a

27" monitor (iiyama, Japan). Participants were recruited among staff and students at the

University of Exeter, aged 18-70 with normal or corrected-to-normal vision and not

knowingly colour-blind. The experiments complied with the declaration of Helsinki and

received ethical approval from the University of Exeter's CLES Cornwall Ethics committee

319   (application ID: 512348, date approved: 22/03/22). Participants received an information

320   sheet presenting the rationale and rules of the search task, and signed a consent form to

321   approve the use of their anonymised data in this study.

323   Results

Case study - pattern evolution in field trials

Prey populations showed evidence of improved camouflage across generations, though

results varied between background types and experimental runs (Fig 4). In the first run,

colour and luminance differences between prey patterns and the background decreased, as

expected, for the population shown on smooth gravel, but actually increased slightly for the

population on furrowed gravel (linear model (lm), generation:background, $F_{2,570}$ = 7.625,

$F_{2,570}$ = 17.425, p < 0.001, for colour difference between means and the weighted average

colour difference respectively; $F_{2,570}$ = 18.997, p < 0.001 for luminance; Fig 4a(i-iii)).

Conversely, pattern differences decreased across generations in the more complex

furrowed treatment, but increased in the smooth (lm, generation:background, $F_{1,572}$ =

16.282, p < 0.001; Fig 4a(iv)), suggesting that the relative importance of pattern and colour

matching varied between backgrounds. In the second run, luminance difference decreased

in both populations, but most strongly on the furrowed backgrounds, (lm,

generation:background, $F_{1,812}$ = 6.869, p = 0.009; Fig 4b(iii)). However, there was little

improvement in colour-matching: differences between mean target and background colours

declined across generations only on the furrowed background (lm, generation:background,

$F_{2,810}$ = 10.761, p < 0.001; Fig 4b(i)) while the weighted average measure found little change

in that population, and an increase in colour difference across generations on the smooth

background (lm, generation:background, $F_{1,812}$ = 28.069, p < 0.001; Fig 4b(ii)). Unlike in the

first run, there were no significant trends in pattern difference (lm, generation:background,

$F_{2,810}$ = 1.328, p = 0.266; background, $F_{1,812}$ = 1.524, p = 0.217; generation, $F_{2,813}$ = 0.975, p =

0.378; Fig 4b(iv)). When improvements in background-matching were observed, effect sizes

were small, and targets remained conspicuous, with colour and luminance differences well

above accepted detection thresholds of 1 to 3 ΔS or ΔL. For example, in the smooth

background population in the first run, colour differences between means declined from

10.28 ΔS in generation 0 to 9.32 ΔS in generation 11 (Fig 4a(i)). Luminance differences

reached lower levels, suggesting that luminance matching was more critical for camouflage,

but this could be due to lower starting values; for example, in the same population, mean

luminance difference started at 7.07 ΔL in generation 0, reaching 3.47 ΔL by generation 11

(Fig 4a(iii)).

Fig 4: Metrics of camouflage efficacy across generations, in targets from the first (a) and

second run (b) of field trials, alongside expected trends for improved camouflage (c). All

plots display targets shown against smooth backgrounds in the left panel, and furrowed

backgrounds on the right. Trend lines and shaded ribbons represent final model predictions,

with 95% confidence intervals. Generation was always initially fitted as a second-degree

polynomial; non-linear trends are shown if the polynomial term was significant in final

models.

Most consistently, targets in all populations displayed increased edge disruption across

generations (Fig 4a,b(v)); this improvement appeared to level off in the population shown

on furrowed backgrounds in the first run (lm, generation:background type, $F_{2,570}$ = 4.754, p =

0.009), but not in the second (lm, generation:background, $F_{2,810}$ = 1.976, p = 0.139;

background, $F_{1,812}$ = 0.531, p = 0.467; generation, $F_{1,814}$ = 72.172, p < 0.001).

Evolution in screen-based replicates

In the screen-based experiments, camouflage metrics reveal clear differences between

populations of prey utilising the full and narrow colour spaces. For populations using the

narrow colour space, patterns improved in every camouflage metric across generations (Fig

5a). There were significant decreases in luminance, colour and pattern differences between

targets and the backgrounds (lme, for luminance: generation, χ^2_1 = 114.07, p < 0.001; for

colour: generation, χ^2_2 = 191.31, p < 0.001; for pattern: generation, χ^2_1 = 8.466, p = 0.004),

and an increase in edge disruption in target patterns (lme, for GabRat in L: generation, χ^2_1 =

8.043, p = 0.005), all with no significant differences between populations shown against

smooth or furrowed backgrounds (see S1 Table). For the weighted average colour difference

metric, there was a significant effect of the interaction between generation and background

type, describing different patterns of overall decrease in colour differences

(generation:background type, χ^2_2 = 7.434, p = 0.024; Fig 5a(iii)). These improvements in

target camouflage made it harder for participants to find them, as suggested by a significant

increase in the time taken to capture the first target on each slide over successive

generations for the narrow colour space, regardless of background type (lme, generation:

background, $\chi^2_2 = 4.762$, p = 0.093, background: $\chi^2_1 = 3.826$, p = 0.050, generation: $\chi^2_2 = 6.340$,

p = 0.012 ; Fig 5a(vi)).

By contrast, for the full colour space, there was substantial variation in the trajectory of

pattern evolution between replicate populations, making it difficult to see consistent

evidence of selection for improved camouflage (Fig 5b). There were no significant trends in

luminance or pattern difference between the targets and the background across

generations and no evidence of increased edge disruption in prey patterns (Fig 5b(i), (iv-v),

S2 Table). There was however some evidence for improved colour-matching overall:

average colour difference between targets and backgrounds decreased slightly across

generations (lme, generation, $\chi^2_2 = 7.256$, p = 0.027), and using the weighted average colour

difference metric suggests that this improvement was greater against the smooth

backgrounds (lme, generation:background, $\chi^2_2 = 6.965$, p = 0.031, Fig 5b(ii-iii), S2 Table).

Capture time was shaped by the interaction between generation and background type, but

did not increase overall (lme, generation: background, $\chi^2_2 = 10.832$, p = 0.004; Fig 5b(vi)).

Fig 5: Changes in camouflage metrics in the screen-based experiments, for prey

populations using the narrow (a) and full (b) colour spaces. Plots show the mean and

standard error of each metric per generation, along with predictions from the best statistical

models. Shades of each colour represent different populations, with filled dots for those

shown on smooth backgrounds, and open dots for furrowed backgrounds. Dotted lines

represent population-level predictions, and thicker lines overall trends for fixed effects only.

Results from the control populations, simulating the effects of neutral drift, confirm that

consistent selection for improved camouflage was only seen in experimental populations

using the narrow colour space. Under those conditions, there were significant differences

between the trajectories of experimental populations, selected by volunteers, and control

replicates across generations: average colour and luminance differences between target

patterns and their backgrounds decreased in the selected populations, while they increased

slightly in the controls (lme, generation:scenario, for luminance χ^2_1 = 9.443, p = 0.002, for

colour, χ^2_1 = 15.557, p < 0.001; S6 Fig). Similar trends for the weighted average colour

difference metrics and edge disruption were not significant, suggesting that volunteers in

our experiment were selecting primarily for improved colour and luminance match (see S6

Fig, S3 Table). However, for the full colour space, analysing all replicates together revealed

no significant interaction between scenarios (control versus experimental) and generation,

for any metrics (see S6 Fig, S4 Table).

Discussion

This study presents methods to support investigations of camouflage evolution using

genetic algorithms in the field, with prey items exposed to wild predators. Our detailed case

study describes how to implement these tools, from modifications of the genetic algorithm

software to techniques for making artificial prey and recording predation events, and

demonstrates the feasibility of this approach. Free-flying birds reliably interacted with

multiple generations of prey, and prey patterns improved in several metrics of camouflage

efficacy, suggesting the potential for adaptive evolution.

However, this case study also highlights important factors to consider when designing this

type of evolutionary experiment. Trends for improved camouflage in the field were

relatively weak, and inconsistent among runs with different predators and background

types. We tested whether these findings were linked to the starting colour space available

for prey patterns, using screen-based search tasks with human volunteers, which closely

replicated the field trials and were much quicker to run. These experiments revealed

substantial variation in outcomes produced by individual participants, when target patterns

utilised the same colour space as in the field: while camouflage improved across generations

in some populations of prey, others became more conspicuous, such that few overall trends

for improved camouflage were detected, and populations were not significantly different

from control replicates, simulating neutral drift. These results were unexpected, as similar

psychophysics experiments in artificial evolution typically find much clearer trends towards

improved camouflage [12]. However, the number of prey and the way they are presented

strongly affect learning in this type of experiment [33]. Here, the presence of multiple prey

per image increases the influence of random factors, such as the position of prey relative to

one another. Under these circumstances, participant learning outstripped target evolution,

highlighting the unexplored power of predator learning in holding back the evolution of

camouflage from a conspicuous evolutionary start point, a scenario that may be relevant to

aposematic species, for example. Predator cognition, encompassing learning, neophobia,

forgetting, recognition errors and adaptive decision-making, is a key consideration in the

evolution and maintenance of aposematism as a signalling strategy [34,35]. Theoretical

models suggest that elements of both predator psychology and prey ecology, such as prey

density and aggregation, can contribute to the initial evolution of aposematism [36], and

our set-up presents opportunities to investigate these effects, by varying the profitability

and arrangement of prey in both the field and online systems. Predator behaviour and

cognition are similarly important for the evolution of camouflage yet have received less

attention in this context [37], so further experiments with real predators could also shed

more light on the conditions required for crypsis to evolve, given different predator

communities. In our experiments, the screen-based trials suggest the trajectory of evolution

is essentially random when prey utilise the full colour space, so any specific results from the

field case study should be interpreted with caution.

By contrast, the screen-based experiments showed that populations of prey utilising a

narrower colour space, based on colours found in the backgrounds, were not only more

cryptic from the start, but also consistently evolved to be better camouflaged over

successive generations, and clearly behaved differently from control replicates. We

therefore predict that field trials would similarly yield greater improvement in camouflage if

the prey patterns started with a more restricted colour space, based on the colours of the

background. In addition, very few metrics of camouflage efficacy revealed differences in

prey pattern evolution between populations displayed against the smooth and furrowed

backgrounds, even in the screen-based experiments with the narrow colour space. In the

field runs, trends did vary between background types, but there was little consistency across

metrics, or between repeats, and thus no clear sense that selection was stronger for

populations on the simpler smooth background. This suggests that the variation in three-

dimensional complexity between these backgrounds was also insufficient to select for

different evolutionary trajectories in this scenario. As such, these experiments represent a

proof-of-principle for virtual evolution in the wild, but future applications should carefully

consider starting parameters, to better calibrate the difficulty of the task.

Researchers attempting similar work would benefit from testing their starting parameters in

pilot experiments with human volunteers as "predators", to establish conditions in which

robust selection can be expected to take place before running more time-consuming

experiments in the field. To that end, the methods described here present a significant

advantage, as CamoEvo's flexibility allows users to set up screen-based experiments that

closely mirror their field setups, and rapidly collect data with human volunteers. Equally, the

limitations of our field experiments highlight the difficulty of predicting the conditions under

which prey patterns can be expected to evolve, which is in of itself an interesting area to

explore. Using these field methods to investigate the scenarios that favour the evolution of

camouflage in response to real predators would be extremely valuable, to better

understand the role of predation as a selective force. In particular, the impact of the starting

colour space available to prey has received little attention, yet is likely to be a significant

constraint for the subsequent evolution of prey patterns. Beyond prey appearance, initial

conditions are an important determinant of evolutionary trajectories more generally. For

example, certain traits can promote or restrict the potential for adaptive radiation (eg. when

competition between small individuals for a shared resource limits the emergence of larger

individuals that could specialise on alternative resources [38]) while more evolutionarily

labile traits can facilitate the spread of invasive species, or boost the resilience of native

species they encounter [39]. So-called "historical difference experiments", testing how

populations that previously evolved under different conditions perform in a new situation,

provide a means to explore the role of starting conditions and historical contingencies in

evolution [40]. Our set-up provides opportunities to run this kind of experiment, potentially

by combining screen and field experiments for efficiency, and approach these broad

evolutionary questions through the study of pattern evolution.

The system we present here is designed to encourage future research on prey appearance,

by providing open-access tools for every step of the process, from pattern generation and

the genetic algorithm framework [12], to implementation in the field, and image analysis

software for quantifying changes in prey properties [19,20]. Within CamoEvo and

CamoPrint, parameters such as target shape, colour space and genetic algorithm rules are

fully customisable, while the field set-up can be modified to meet individual needs, for

different predators and backgrounds. This should facilitate the application of virtual

evolution to investigating a wide range of visual pattern types and functions. In particular,

identifying optimal camouflage patterns remains a key question, with many practical

applications. Recent studies tackling this problem have gained useful insights by defining

optimal patterns based on the statistical properties of the background [41], or using deep

learning to extrapolate from human responses to a limited set of phenotypes [3]. Yet

genetic algorithms present an especially powerful tool for studying camouflage

optimisation, particularly when paired with machine learning techniques [10,11]. Crucially,

they account for the effects of repeated interactions between predators and prey, though

implementation in screen-based experiments restricts their interpretation. Associating

these tools with predation experiments in the field provides opportunities to investigate

optimal camouflage while simultaneously considering a wider range of influential factors,

from background properties and viewing conditions to predator vision, learning and prey

population dynamics. Beyond camouflage, field-based evolutionary approaches supported

by these tools could also provide valuable new insights into other forms of anti-predator

colouration, such as aposematism or distant-dependent signalling.

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Conflict of Interest Statement

The authors declare no conflict of interest.

Author Contributions

JT: Conceptualization, Methodology, Supervision, Writing – review & editing

GRAH: Conceptualization, Methodology, Investigation, Visualization, Writing – review &

editing

ESB: Methodology, Investigation, Formal Analysis, Visualization, Writing – original draft,

Writing – review & editing

Data Availability

Data and analysis code are available as supplementary material; code for tools used in this

paper can be found on GitHub:

https://github.com/GeorgeHancock471/CamoWild_Repository_CE.v1.2

The tools used for creating printed targets, including site randomisation have been

condensed and modified for greater ease of use in CamoEvo Version 2.0 and will be

included in all future versions. Instructions on using these tools are provided in the

CamoEvo 2.0 handbook: https://github.com/GeorgeHancock471/CamoEvo-v2.0-

2022_Plugins/tree/main

551 References

1. Stevens M, Merilaita S. Animal camouflage: current issues and new perspectives. Philos Trans
R Soc London B Biol Sci. 2009;364(1516):423–7.

2. Kjernsmo K, Whitney HM, Scott-Samuel NE, Hall JR, Knowles H, Talas L, et al. Iridescence as
camouflage. Curr Biol. 2020;30(3):551–5.

3. Fennell JG, Talas L, Baddeley RJ, Cuthill IC. Optimizing colour for camouflage and visibility
using deep learning: the effects of the environment and the observer's visual system. J R Soc
Interface. 2019;16:20190183.

4. Bond AB, Kamil AC. Visual predators select for crypticity and polymorphism in virtual prey.
Nature. 2002;415:609–13.

5. Troscianko J, Nokelainen O, Skelhorn J, Stevens M. Variable crab camouflage patterns defeat
search image formation. Commun Biol. 2021;4(1):1–9.

6. Whitley D. A genetic algorithm tutorial. Stat Comput. 1994;4(2):65–85.

7. Bond AB, Kamil AC. Spatial heterogeneity, predator cognition, and the evolution of color
polymorphism in virtual prey. Proc Natl Acad Sci. 2006;103(9):3214–9.

8. Sherratt TN, Pollitt D, Wilkinson DM. The evolution of crypsis in replicating populations of
web-based prey. Oikos. 2007;116:449–60.

9. Reynolds C. Interactive evolution of camouflage. Artif Life. 2011 Apr 1;17(2):123–36.

10. Talas L, Fennell JG, Kjernsmo K, Cuthill IC, Scott-Samuel NE, Baddeley RJ. CamoGAN: evolving
optimum camouflage with Generative Adversarial Networks. Methods Ecol Evol.
2020;11(2):240–7.

11. Fennell JG, Talas L, Baddeley RJ, Cuthill IC, Scott-Samuel NE. The Camouflage Machine :
optimizing protective coloration using deep learning with genetic algorithms. Evolution.
2021;1–11.

12. Hancock GRA, Troscianko J. CamoEvo: an open access toolbox for artificial camouflage
evolution experiments. Evolution. 2022;870–82.

13. Sherratt TN, Beatty CD. The evolution of warning signals as reliable indicators of prey
defense. Am Nat. 2003;162(4):377–89.

14. Hughes AE, Griffiths D, Troscianko J, Kelley LA. The evolution of patterning during movement

in a large-scale citizen science game. Proc R Soc B Biol Sci. 2021;288(1942).

15.   Lawson DA, Rands SA. The evolution of floral guides: Using a genetic algorithm to investigate
582   the evolution of floral cue arrangements. Biol J Linn Soc. 2018;123(4):739–53.

583   16.   Allen JA, Cooper JM, Hall GJ, McHenry C. 'Evolving pastry': a method for simulating
584   microevolution. J Biol Educ. 1993;27(4):274–82.

585   17.   Cooper JM, Allen JA. Selection by wild birds on artificial dimorphic prey on varied
586   backgrounds. Biol J Linn Soc. 1994 Apr 1;51(4):433–46.

587   18.   Orteu A, Jiggins CD. The genomics of coloration provides insights into adaptive evolution. Nat
588   Rev Genet. 2020;21(8):461–75.

589   19.   Troscianko J, Stevens M. Image calibration and analysis toolbox - a free software suite for
590   objectively measuring reflectance, colour and pattern. Methods Ecol Evol. 2015;6(11):1320–
591   31.

592   20.   van den Berg CP, Troscianko J, Marshall NJ, Cheney KL, Endler JA. Quantitative Colour Pattern
593   Analysis (QCPA): a comprehensive framework for the analysis of colour patterns in nature.
594   Methods Ecol Evol. 2020;11:316–32.

595   21.   Merilaita S. Visual background complexity facilitates the evolution of camouflage. Evolution.
596   2003;57(6):1248–54.

597   22.   Xiao F, Cuthill IC. Background complexity and the detectability of camouflaged targets by
598   birds and humans. Proc R Soc B Biol Sci. 2016; 283:20161527.

599   23.   Murali G, Mallick S, Kodandaramaiah U. Background complexity and optimal background
600   matching camouflage. Behav Ecol Sociobiol. 2021;75(4).

601   24.   Rowe ZW, Austin DJD, Chippington N, Flynn W, Starkey F, Wightman EJ, et al. Background
602   complexity can mitigate poor camouflage. Proc R Soc B Biol Sci. 2021;288(1963).

603   25.   Lovell PG, Tolhurst DJ, Párraga CA, Troscianko J, Troscianko T, Baddeley R, et al. Stability of
604   the color-opponent signals under changes of illuminant in natural scenes. J Opt Soc Am A.
605   2005;22(10):2060.

606   26.   Troscianko T, Benton CP, Lovell PG, Tolhurst DJ, Pizlo Z. Camouflage and visual perception.
607   Philos Trans R Soc B Biol Sci. 2009;364(1516):449–61.

608   27.   Schneider CA, Rasband WS, Eliceiri KW. NIH Image to ImageJ: 25 years of image analysis. Nat
609   Methods. 2012;9(7):671–5.

610   28.   Vorobyev M, Osorio D. Receptor noise as a determinant of colour thresholds. Proc R Soc B
611   Biol Sci. 1998;265:351–8.

612   29.   Renoult JP, Kelber A, Schaefer HM. Colour spaces in ecology and evolutionary biology.
613   2017;92:292–315.

614   30.   Spottiswoode CN, Stevens M. Visual modeling shows that avian host parents use multiple
615   visual cues in rejecting parasitic eggs. Proc Natl Acad Sci. 2010 May 11;107(19):8672–6.

616   31.   Troscianko J, Skelhorn J, Stevens M. Quantifying camouflage: how to predict detectability
617   from appearance. BMC Evol Biol. 2017;17(1):7.

618   32.   R Development Core Team. R: a language and environment for statistical computing. Team
619   RDC, editor. R Foundation for Statistical Computing. 2015. Available from: http://www.r-

project.org

33. Troscianko J, Lown AE, Hughes AE, Stevens M. Defeating crypsis: detection and learning of
camouflage strategies. PLoS One. 2013;8(9):e73733.

34. Servedio MR. The effects of predator learning, forgetting, and recognition errors on the
evolution of warning coloration. Evolution. 2000;54(3):751–63.

35. Skelhorn J, Halpin CG, Rowe C. Learning about aposematic prey. Behav Ecol. 2016;27(4):955–
64.

36. Speed MP. Can receiver psychology explain the evolution of aposematism? Anim Behav.
2001;61(1):205–16.

37. Skelhorn J, Rowe C. Cognition and the evolution of camouflage. Proc R Soc B Biol Sci.
2016;283(1825):20152890.

38. Ten Brink H, Seehausen O. Competition among small individuals hinders adaptive radiation
despite ecological opportunity. Proc R Soc B Biol Sci. 2022;289(1971).

39. Perkins TA. Evolutionarily labile species interactions and spatial spread of invasive species.
Am Nat. 2012;179(2):E37–54.

40. Blount ZD, Lenski RE, Losos JB. Contingency and determinism in evolution: replaying life's
tape. Science. 2018;362(6415).

41. Michalis C, Scott-Samuel NE, Gibson DP, Cuthill IC, Michalis C. Optimal background matching
camouflage. 2017; 284:20170709.

640 Supporting Information captions

SI1: Additional methodological information for case study field trials

S1 Fig. Evidence of predation from a crow (*C. corone*) in run 1 (a), and a robin (*E. rubecula*), great tit
(*P. major*) and blackbird (*T. merula*) in run 2 (b-d).

S2 Fig. Prey patterns in the field trials: generation 0 (left) and the final generation in runs 1 (middle)
and 2 (right) for populations shown on smooth (top) and furrowed backgrounds (bottom).

S3 Fig. Altered target shading method, showing the base method (top-down) on the left and the
custom method (centre-edge) on the right.

SI 2: Additional information for screen-based search tasks

S4 Fig. Target image modification for screen-based replicate experiments.

S5 Fig. Schematic of the screen-based search tasks, showing the targets in generation 0 and 15, for
populations using the full and narrow colour spaces.

S6 Fig. Changes in camouflage metrics in the screen-based experimental populations and control

replicates, using the narrow (a) and full (b) colour spaces.

S1 Table. Model simplification tables for full models testing changes in metrics of camouflage

efficacy in the screen-based experiment, for the narrow colour space.

S2 Table. Model simplification tables for full models testing changes in metrics of camouflage

efficacy in the screen-based experiment, for the full colour space.

S3 Table. Model simplification tables for full models testing changes in metrics of camouflage

efficacy between control and experimental populations of the screen experiment, for the narrow

colour space.

S4 Table. Model simplification tables for full models testing changes in metrics of camouflage

efficacy between control and experimental populations of the screen experiment, for the full colour

space.

S1 Video. Example predations in field trials.

Experimental design

Prey shape & size

Prey orientation

Population size

Background type

Prey grouping & placement

Predators & setting

Starting colour space

Implementation

CamoEvo & CamoPrint

Evaluation

Colour & pattern analysis methods in CamoEvo & QCPA/MICA Toolbox (ImageJ), or other software

- Colour, luminance & pattern similarity
- Edge disruption, salience

Figure 1

Figure 2

Figure 3

Figure 4

Figure 5