Light environment influences mating behaviours during the early stages of divergence in tropical butterflies

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ABSTRACT

Species divergence is facilitated when traits under divergent selection also act as mating cues. Fluctuations in sensory conditions can alter signal perception independently of adaptation to contrasting sensory environments, but how this fine scale variation affects behavioural isolation has received less attention, especially in terrestrial organisms. The warning patterns of *Heliconius* butterflies are under selection for aposematism and act as mating cues. Using computer vision, we extracted behavioural data from 1481 hours of video footage for 387 individuals. We show that the putative hybrid species *H. heurippa* and its close relative *H. timareta linaresi* differ in their response to divergent warning patterns, and that these differences are strengthened with increased local illuminance. Trials with live individuals reveal low-level assortative mating that are sufficiently explained by differences in visual attraction. Finally, results from hybrid butterflies are consistent with linkage between a major warning pattern gene and the corresponding behaviour, though the differences in behaviour we observe are unlikely to cause

rapid reproductive isolation as predicted under a model of hybrid trait speciation. Overall, our

results highlight that the role of ecological mating cues for behavioural isolation may depend on

the immediate sensory conditions during which they are displayed to conspecifics.

Keywords: *Heliconius* – Ecological Speciation – Sensory Environment – Magic traits – Hybrid

Trait Speciation – Computer vision

BACKGROUND

During ecological speciation, barriers to gene flow evolve as a result of ecologically-based

divergent selection [1]. These barriers are generally expected to build up gradually over time [2],

with premating isolation evolving early alongside ecological differences [1,3,4]. The evolution of

premating isolation is facilitated when traits under ecological selection act as mating cues

(sometimes known as 'magic traits' [5,6]), as this allows divergent natural selection acting on

ecological traits to be transferred to mating behaviours. However, the strength of behavioural

barriers may be influenced by the immediate, and perhaps rapidly changing, sensory conditions,

but this has received relatively little empirical attention (but see [7–13]). One reason may be that

robustly detecting these effects during the early stages of divergence, when they may be most

relevant, presents a substantial empirical challenge.

It is well established that the sensory environment can alter signal detection and

perception [14,15]. Colour perception depends not only on an object's reflectance spectrum, but

also the illumination, available light spectrum and background, all of which may change within

seconds and over very short distances [16]. This can affect within-population preferences. For

example, red colouration in male *Habronattus* spiders is only an efficient mating signal if

presented in broad-spectrum sunlight [17]. Other sensory modalities can be also affected: Female

swordtail fish are strongly attracted to the chemical signals of conspecific males in clean water,

but not in polluted water [7]. Similarly, urban noise can disrupt the transmission of bird songs

and interfere with acoustic-based mate choice [9].

The sensory environment may affect the evolution of reproductive isolation in two main

ways. First, adaptations to meet the ecological needs of different sensory environments can

influence female preferences, subsequently driving divergence in male signals, and leading to

reproductive isolation through sensory drive [18]. For example, in two closely related *Pundamilia*

cichlid fishes, adaptation of the visual system to the local environment is associated with

divergent female mate preference for male colouration [19]. Second, prevailing environmental

conditions may alter the efficacy of signals used in mate choice [14], so that individual mating

preferences may act as an important reproductive barrier under some sensory conditions, but not

others ('context-sensitive preferences' [20]). If the strength of preferences depends on the sensory

environment, then this will influence their contribution to reproductive isolation. However,

compared to the role of sensory adaptation, the immediate influence of local sensory conditions

on mating behaviours, and how this may relate to the evolution and maintenance of new species,

has been less-studied, especially in terrestrial organisms.

By affecting context-sensitive preferences, the sensory conditions during signalling may

also influence the strength of genetic associations (*i.e.* linkage disequilibrium, LD) between

mating and ecological traits, which are typically required for speciation to proceed when gene

flow persists [21]. Specifically, when ecological traits act as mating cues, LD (between

ecological and preference loci) will arise as a natural consequence of mating preferences, but this

will be proportional to the strength of preference [22], which may be influenced by the sensory

environment. Regardless, unless preferences are very strong, more robust coupling between

mating and ecological components of reproductive isolation likely requires genetic architectures

that impede recombination (or one-allele mechanisms, see [23]). These include physical linkage,

which may be further strengthened by genomic rearrangements like inversions, or – in the

extreme – pleiotropy, where distinct traits are controlled by the same allele [24]. To date, studies

have reported physical linkage between behavioural and ecological components of reproductive

isolation for a few animal taxa, including pea aphids [25], fish [26] and butterflies [27].

Heliconius butterflies are known for their bright warning patterns, which are often under

selection for Müllerian mimicry [28]. Closely related *Heliconius* species frequently – but not

always – have very different wing patterns, which additionally act as mating cues (e.g. [29–32]).

This contributes to assortative mating because males almost invariably prefer wing patterns

resembling their own, and warning patterns in *Heliconius* are considered one of the best examples

of 'magic traits' in nature [6,29,33]. Variation in warning pattern is largely controlled by a few

genes; the genetics of the corresponding visual mate preference are less well known, though

recent work implicates a handful of genes associated with neural signalling in tight linkage to the

colour pattern gene *optix* [27,34].

There is substantial evidence that colour pattern alleles have introgressed between

otherwise separately evolving *Heliconius* lineages (e.g. [35]). In particular, taxa within the

heurippa-timareta group, found in the eastern slopes of the South-American Andes, have

acquired red colour pattern elements from local races of *H. melpomene* via introgression of *optix*

alleles [36]. This has frequently led to near perfect mimicry between local races of *H. timareta*

and *H. melpomene*; elsewhere, however, the resulting colour patterns are not shared with local

Heliconius species. In particular, *H. heurippa* has a unique red-yellow banded forewing pattern

(Fig. 1A). Its close relative *H. timareta linaresi*, which is assumed to represent the ancestral wing

colour pattern of the *heurippa-timareta* group [37], only displays a yellow band (Fig. 1A) and

also does not have an obvious co-mimic. These two populations are geographically adjacent (Fig.

1A) and presumably share a contact zone. Despite their nominal status as separate species, *H. t.*

linaresi and *H. heurippa* likely represent an early stage of divergence; hybrids between *H. t.*

linaresi and *H. heurippa* are fully viable and fertile [38] and any post-mating isolation is

probably limited to selection acting against immigrant warning patterns.

The red pattern element of *H. heurippa* is presumably maintained by strong frequency-

dependent selection imposed by predators [39]. However, its effectiveness as intraspecific signal

may depend on how it is perceived by conspecifics under natural sensory conditions. In this way,

premating reproductive barriers may depend on interactions between the signal, environment and

receiver. *H. melpomene* are broadly separated from *H. heurippa-timareta* (along with closely

related *H. cydno*) taxa across a gradient of open to closed forest and decreasing light intensity

[32,40]. These contrasting sensory environments are predicted to alter how colour patterns are

perceived by butterfly visual systems [41]. While *H. heurippa* and *H. t. linaresi* are not

ecologically isolated in this way, their forest habitats are highly heterogenous and *Heliconius* are

known to settle with their wings open in more brightly illuminated 'light patches' [42]. Whether

changes in illuminance affect male behaviours has not been investigated, but such an effect

would suggest that the efficacy of wing patterns as premating reproductive barriers may depend

on fluctuations in sensory conditions as females move through the environment.

To test whether warning patterns contribute to premating isolation between *H. heurippa*

and *H. t. linaresi*, we used a novel computer vision pipeline to extract behavioural data from

video footage, alongside choice trails to determine levels of assortative mating. Our data,

including ~17,000 behavioural interactions for 387 individuals, allowed us to test (i) whether the

two species show differences in visual attraction towards con- and hetero-specific patterns and

(ii) whether these differences segregate with the colour patterns, consistent with physical linkage

between the behavioural and colour patterning genes. Measuring illuminance in real-time at each

female pattern also allowed us to ask (iii) whether these behavioural differences are influenced by

fluctuating local light conditions.

MATERIAL AND METHODS

(a) Butterfly collection, maintenance and crossing design. We maintained outbred stocks of *H.*

timareta linaresi and *H. heurippa*, established from wild individuals (Table S1), between January

2016 and September 2019 at the Universidad del Rosario insectaries in La Vega, Colombia

(4°59' N, 74°20' W, elevation 1257 m). We generated F_1 hybrids and backcross hybrids to each

species (BL, backcross to *H. t. linaresi* and BH, backcross to *H. heurippa*) (Table S2). All

butterflies were supplied with ~10% sugar solution and *Lantana*, *Psiguria* and/or *Gurania spp.*.

Females were kept individually, with *Passiflora* for oviposition, whereas males were kept in

groups. Eggs were collected every other day and the caterpillars were raised until eclosion in

plastic cups, and fed on fresh *Passiflora* leaves.

Figure 1: A) Locations in Colombia where *H. heurippa* (red) and *H. timareta linaresi* (blue) are known to occur. B) Video recording setup. Mounted females were presented simultaneously: Each was filmed and an associated light logger recorded illuminance every second. 'Tripod' was not casting shade on light logger or mounted females at any time. C) *Post-hoc* motion detection pipeline. A difference image is formed between each frame and its predecessor (identifying pixels with a change in value). Thresholding, blurring and again thresholding pick out significant local changes ('signals'). 'Signal' frames (and surrounding frames) were extracted.

(b) Trials with mounted females. To test for differences in visual mating behaviours, we assayed males in choice trials with dead mounted *H. heurippa* and *H. t. linaresi* females presented simultaneously in an exposed 4x4x2m insectary, in which light conditions varied due to changes in cloud coverage and patterns of shade cast by vegetation at certain daytimes. Virgin females were frozen with their wings spread on the day of eclosion and kept at -20°C for >168h. They were then dried and subsequently washed in hexane to remove residual cuticular hydrocarbons and other pheromones, and mounted onto a small piece of wire. Throughout the

experiment, we used 21 *H. heurippa* and 23 *H. t. linaresi* mounted females, randomly choosing a

pair each day. Mounted females were individually attached to a horizontal wire (~70cm above

ground) at one of six locations within the experimental cage. This location was changed every

hour (hereafter 'trial'), and the new location was chosen randomly (but female types were never

in the same location twice during the same day). A *GoPro Hero 5 Black*™ (GoPro, San Matteo,

CA) camera (settings and equipment in Table S3) was installed at each position, and ~50cm

above the respective mounted female (Fig. 1B). At same height as the mounted female at ~30cm

distance, we attached a *HOBO*™ *UA-002-64* logger (sensor facing up) to measure illuminance

[*lux*] every second (Fig. 1B). Cameras and light loggers were synced using *GoPro Quik*™ and

HOBOware™ software, respectively, allowing us to match video frames with light

measurements.

Most of the virgin naive males matured in one separate communal cage before being

introduced into the experimental cage >4 days after eclosion, where they were tested in mixed

groups (median group size = 22). Males were numbered and received a unique code of dots on

the dorsal side of the wings, allowing identification from videos. Each male was tested over

multiple days (median = 12d). We recorded an average of 3.01h of material on each of the 246

recording days; conducting behavioural trials across different seasons and at different daytimes

allowed us to capture a variety of light conditions.

(c) Computer vision and video analyses. We used a custom motion-detection pipeline, which

post-hoc discarded video footage with no activity. The detection of frames with motion

('signals') was based on difference imaging and consecutive steps of blurring and thresholding,

as implemented in the *OpenCV* library available for C++ (Fig. 1C). Not all of the frames of male

motion sequences made it over the threshold, so we determined the 'signal'-frames and the

surrounding frames one sec before and after a signal in R [43]. Reduced videos were created with

the *OpenCV* library in C++. (Code is accessible at:

github.com/SpeciationBehaviour/visual_preference_heurippa_linaresi). Video material was

processed on an HP^{TM} Desktop computer (i7-7700 CPU, 4 cores), at runtimes ~80% of video

duration. All videos were processed under the same threshold (145) and blur (30) settings.

Remaining footage was curated manually at 66.6% speed using the $MPCHC^{TM}$ *player*. We

recorded three behaviours: 'approach' (male is changing its flight direction towards the mounted

female, resulting in a curve or circling motion), 'courtship' (sustained hovering above mounted

female) and 'sitting' (male sits down on mounted female). Behaviour types were combined for

subsequent analyses.

(d) Tetrad experiments with live females. We performed 'tetrad' trials with virgin males and

females to test for assortative mating. For each trial, sexually mature *H. heurippa* and *H. t.*

linaresi males (one of each) were allowed to acclimatize for 15 mins in a 2x4x2m insectary, at

which point *H. heurippa* and *H. t. linaresi* virgin females (one of each) were introduced. Once the

first mating occurred, the experiment was stopped.

(e) Statistical analyses. We measured illuminance at both mounted females for 83% of recorded

behaviours. Illuminance, measured in *lux*, is the intensity of light falling onto a surface. We

log10-transformed *lux* measures (log-illuminance) and scaled (and centred) each set of log-

illuminance measurements, making the choice of logarithm base irrelevant for the analyses.

Analyses were conducted in R [43] (supplemental R Markdown and

https://github.com/SpeciationBehaviour/visual_preference_heurippa_linaresi). Posteriors will be

described with a 95% equal-tailed credible interval and the mean as point estimate. We analysed

data from trials with mounted females using logistic regression with the *brms* package [44], an

interface to the Bayesian software Stan [45]. Male behaviours directed towards the *H. t. linaresi*

or *H. heurippa* mounted females were fitted as binary Bernoulli-distributed response variable (*i.e.*

0 and 1, respectively); estimates from the model can be understood as a proportion of interactions

with the mounted *H. heurippa* female, with higher values indicating stronger relative attraction to

the *H. heurippa* 'female' and lower values indicating stronger relative attraction to the *H. t.*

linaresi 'female'. Models initially included all possible nested variations of the fully saturated

model explaining effects of 1) male type, 2) log-illuminance at the *H. heurippa* 'female', 3) log-

illuminance at the *H. t. linaresi* 'female', and their interactions. Individual ID and trial were fitted

as random effects.

To test for 'species' differences, we initially fitted models for the 'pure' males (male type

= '*H. heurippa*' or '*H. t. linaresi*'). Segregation of the red bar in BL hybrids (controlled by alleles

at *optix* [46]) allowed us to test for linkage between colour and preference loci (see [27,47]) (here

male type = 'red' (*Bb* genotype) or 'no red' (*bb* genotype) and brood was additionally fitted as a

random effect). To determine which terms to retain [48], we calculated the widely applicable

information criterion (WAIC) and the leave-one-out information criterion (LOOIC) for each set

of models, using the *loo* package [49], and WAIC weights using the *brms* package [44]. Weakly

informative priors (centred around the value for no preference) were set for coefficients

corresponding to the different male types, which gave small prior probabilities for extreme values

very close to 0 or 1. For all other coefficients, default (non-informative) priors were applied. We

also fitted 'categorical illuminance' models adopting the best fitting model structure determined

for each dataset, where values =<median were 'poorly lit', and values >median were 'brightly lit'

(Fig. S1). Posteriors of the estimated marginal means (EMMs) were calculated using the

emmeans package [50]. From this, we retrieved posteriors of contrasts and calculated the

posterior probability (PP) that EMMs differ.

For the tetrad data, we fitted observed counts of each mating outcome (type of male and

female involved) as Poisson-distributed response variable. We included the specific male-female

combination of the mating outcome as fixed factor. Transforming the resulting model estimates

into proportions effectively makes this a multinomial model [51]. Non-informative default priors

were applied throughout. PPs were calculated with the *brms* package [44]. We compared the

observed frequency of each mating combination to the predicted frequencies based on our

measurements of visual preference from the mounted female experiment. Predictions were

derived by multiplying the frequency that a male type was involved in *any* mating combination

by its respective EMMs from the models fitted to the mounted female data. Predictions were

based either on the overall EMM for each type, or the interaction term EMMs from the

categorical model. Posterior distributions for each prediction were calculated using the *binom*

package under the default prior [52].

RESULTS

(a) Species comparisons: i) Divergent visual attraction behaviours in *H. timareta linaresi*

and *H. heurippa* males are dependent on the light environment.

Over 1.5 years we collected 1481 hours of footage. Our computer vision pipeline reduced this to

66 hours requiring human curation (*i.e.* 4.5% of the total footage recorded), including 16,995

behavioural 'interactions' from 387 males (~43.9 per male). These data allowed us to determine

the effects of male type and illuminance on relative visual attraction to the *H. heurippa* mounted

'female' (hereafter 'preference'). The best fitting model for the pure *H. heurippa* and *H. t.*

linaresi males retained male type, log-illuminance at the *H. heurippa* 'female' and their

interaction (Table S4: model #1). Overall, our results suggest that the local light environment

influences the strength of visual attraction.

Across the entire dataset, illuminance at the mounted *H. heurippa* female increased the

difference in preference between the male types, as evidenced by the interaction between male

type and log-illuminance at the *H. heurippa* 'female' (Fig. 2A & S3, PP simple slope *H. heurippa*

> *H. t. linaresi* = 0.996). This was largely driven by an increase in log-illuminance at the *H.*

heurippa 'female' leading to a stronger conspecific preference in *H. heurippa* males (PP simple

slope > 0 = 0.993); there was only limited support for an effect on *H. t. linaresi* males (PP simple

slope < 0 = 0.768). Overall, *H. heurippa* males showed a higher proportion of interactions with

the *H. heurippa* pattern than *H. t. linaresi* males. Although supported with high credibility (PP

relative visual attraction *H. heurippa* > *H. t. linaresi* = 0.969), this difference in preference was

relatively small (0.07, CrI: 0.00 - 0.14) and characterised by considerable within-population

variation (Fig. 2B). Nevertheless, this effect nearly doubled when the *H. heurippa* 'female' was

in brighter light (0.13, CrI: 0.05 - 0.22; Fig 2D) and was absent when the *H. heurippa* 'female'

was poorly lit (0.01, CrI: -0.07 - 0.09; Fig 2C).

Figure 2: The effect of illuminance on relative attraction towards *H. heurippa* 'females' (*i.e.* proportion of

interactions with the *H. heurippa* 'female' as opposed to the *H. t. linaresi* 'female'). **A**: Relative attraction towards

H. heurippa 'females' of *H. t. linaresi* (blue) and *H. heurippa* males (red) under changing illuminance at the *H.*

heurippa 'female'. Illuminance on the x-axis is log-scaled. Coloured area around each regression line represents the

95% credible interval (CrI). Dashed vertical line represents the median log-illuminance used as a cutoff to define the

poorly and brightly lit conditions (below). Relative attraction towards *H. heurippa* 'females' for *H. t. linaresi* males

(blue) and *H. heurippa* males (red): (**B**) across light environments; (**C**) for poorly lit *H. heurippa* 'female'; (**D**) for

brightly lit *H. heurippa* 'female'. Gardner-Altman plots in B-D show the difference between the two male types:

Horizontal lines project from the means of the posteriors for each male type (means and CrIs in Table S5). The mean

and the 95% CrI for the posterior of the difference between the male types are shown on the right. Each point

represents a single individual and its size is scaled to the number of observations. Custom swarmplot was used to

distribute the dots horizontally.

(b) Species comparisons: ii) *H. heurippa* males mate more often with conspecific females in

choice trials.

To test for assortative mating we also conducted 'tetrad' mate choice trials between *H. heurippa*

and *H. t. linaresi*. During 89 tetrad trials we observed 50 con- and 39 heterospecific matings (PP

for positive assortative mating = 0.88). This trend was driven by a higher proportion of

conspecific matings involving *H. heurippa* males (0.405, CrI: 0.305 - 0.508) than heterospecific

matings involving *H. heurippa* males (0.270, CrI: 0.183 - 0.366]) (PP = 0.94). *H. t. linaresi* males

did not participate more frequently in con- rather than heterospecific matings (PP = 0.43) (Fig. 3).

In general, *H. heurippa* males mated more often than *H. t. linaresi* males (60 vs. 29 times). Our

results closely match predictions derived from the mounted females experiment with brightly lit

H. heurippa 'female' (horizontal purple bars in Fig. 3), and, to some extent, those derived from

all of the illuminance conditions combined (Table S6).

Figure 3: *H. heurippa* males show a preference for live, conspecific females in the tetrad experiments. Dashed
vertical line indicates expected proportion under no assortative mating. Posterior distributions for each mating pair
combination are displayed as histograms (red = 95% CrI, orange line = mean). Predictions based on the visual
attraction data from the mounted females experiment with brightly lit *H. heurippa* 'female' are displayed with their
95% CrI as horizontal purple lines.

(c) Hybrid comparisons: Patterns of behaviour in backcross hybrids are consistent with

linkage between colour and preference loci.

Estimates of preference suggest that first generation hybrid (F$_1$) males behave like *H. heurippa*

males (Fig. S4D), perhaps suggesting that *H. heurippa* alleles for attraction to red are dominant;

however, challenging this, BH males seem to display a preference similar to *H. t. linaresi* males

(Fig. S4E). For BL males, the model including all possible interactions was the best fit (Table S7,

model #1). As for the 'pure' males, the six best fitting models included the interaction between

type (*i.e.* red (*Bb*) and non-red (*bb*) wing pattern) and log-illuminance at the *H. heurippa* 'female'

(91% of cumulative WAIC weight for an effect of this interaction term).

As for males from the parental populations, the difference between the two genotypes was

higher when the *H. heurippa* 'female' was in bright light (Fig. 4CD). There was a small effect of

the three-way interaction with illuminance measures at the *H. t. linaresi* 'female' - the difference

was slightly higher when only the *H. heurippa* mounted female was brightly lit (Fig. 4C, 0.11,

CrI: 0.02 - 0.21), as opposed to when both were (Fig. 4D, 0.07, CrI: 0.00 - 0.13). A brightly lit *H.*

t. linaresi mounted female led to a slightly reversed pattern (*i.e. bb > Bb*), but only when the *H.*

heurippa mounted female was poorly lit (Fig. 4B, -0.03, CrI: -0.13 - 0.07). If both mounted

females where poorly lit, there was no difference in preference (Fig. 4A, 0.01, CrI: -0.05 - 0.08).

Across the entire dataset, illuminance at the mounted *H. heurippa* female increased the

differences in preference between the two BL genotypes (Fig. S2B, PP simple slope *Bb > bb* =

0.999). While the slope for *Bb* males is only slightly positive (PP = 0.859), the slope for *bb* males

is strongly negative (PP = 0.999). In contrast to the pure males, we also observed an effect of

illuminance at the *H. t. linaresi* 'female'. Specifically, *bb* males showed a stronger preference for

the *H. heurippa* 'female' when the *H. t. linaresi* 'female' was bright (Fig. S2D & S3, PP =

0.999). Overall, *Bb* males were more likely to interact with the *H. heurippa* 'female' than *bb*

males (Fig. S5). This difference was supported with moderately high probability (PP relative

visual attraction *Bb > bb* = 0.897). Although these represent small effects in absolute terms (0.03,

CrI: -0.02 - 0.08]), this difference accounts for ~50% of the difference in means of the parental

populations, consistent with linkage between colour and preference loci.

Figure 4: The effect of illuminance on relative attraction towards *H. heurippa* 'females' (i.e. the proportion of interactions with the *H. heurippa* 'female' as opposed to the *H. t. linaresi* 'female') for BL hybrid males (no red/*bb* = blue dots; red/*Bb* = red dots): (**A**) with poorly lit *H. t. linaresi* and *H. heurippa* 'females'; (**B**) brightly lit *H. t. linaresi* and poorly lit *H. heurippa* 'females'; (**C**) poorly lit *H. t. linaresi* and brightly lit *H. heurippa* 'females'; and (**D**) brightly-lit *H. t. linaresi* and *H. heurippa* 'females' (means and CrIs in Table S5). Gardner-Altman plots are as in Figures 1B-D.

DISCUSSION

By collecting ~1500h of mate choice data we have shown that the light environment can

influence visual mating behaviours during the early stages of divergence in *Heliconius* butterflies.

Although our data are characterised by considerable individual variation, we observed significant

differences in behaviours of *H. heurippa* and *H. timareta linaresi* males, and this difference is

stronger when the female patterns are more brightly lit. Experiments with live males and females

revealed a degree of assortative mating, and the differences in visual attraction behaviours that

we observe are sufficient to explain this. We have also shown that differences in visual attraction

are associated with the presence of the red forewing band in interspecific hybrids under bright

light conditions, perhaps suggesting physical linkage between an ecologically relevant colour

pattern gene and those for the corresponding behaviour.

Studies of speciation often focus on already diverged groups, which are maintained by

multiple reproductive barriers [53,54], making it difficult to draw conclusions about the role of

the individual barriers and their ecological context. At the other extreme, barriers acting at the

early stages of divergence may be of small effect and, especially in the case of behavioural

phenotypes, require very large datasets to draw robust conclusions, which may not always be

feasible. Behavioural researchers increasingly use computer vision [55], and software for

automated tracking and individual identification is available (e.g. [56,57]). However, applying

these techniques to footage with heterogenous backgrounds, variable light environments and

arenas larger than the camera's field of view remains a challenge. To overcome these limitations,

we combined computer vision, allowing *post-hoc* motion detection, with subsequent human

curation. This permitted a simple low-cost solution, and importantly allowed us to capture frames

before and after motion is detected from many hundreds of hours of video footage.

This large volume of data revealed that increased illuminance at the red mounted *H.*

heurippa female increased the frequency of interactions between *H. heurippa* males with *H.*

heurippa females (relative to *H. t. linaresi* females). *H. heurippa* males were no more likely than

H. t. linaresi males to interact with the *H. heurippa* female when she was poorly lit (lower 50%

quantile of illuminance values; Fig. 2C), but were 1.3 times (*i.e.* 0.57/0.44, Table S5) more likely

to interact with the *H. heurippa* female in brighter light (upper 50% quantile of illuminance

values; Fig. 2D). Similar effects of the light environment have been observed in aquatic

organisms. For example, the attractiveness of red colour during courtship is dependent on the

spectrum of available light, influenced by water depth and turbidity, and recognition of con-

specifics may fail entirely under certain light environments (e.g. [10–12,19]). However, far fewer

studies have directly tested how changing light environments influence mating preference

behaviours that contribute to population divergence in terrestrial organisms (but see [17,58,59]).

The mechanism underlying the differences in behaviour we observe under contrasting

lighting conditions isn't immediately clear. Insects, including butterflies [60], have frequently

evolved colour constancy across light environments [61,62], and although this may only partially

succeed [60,61], we generally expect individuals to be able to distinguish *H. heurippa* and *H. t.*

linaresi patterns under different brightness conditions. Alternatively, male attraction to female

colour patterns might to some degree be 'wavelength-specific', where triggering of behaviours

depends greatly on an object's emission of specific wavelengths and their intensity [62]. Under

this scenario, a colour cue might only trigger a behaviour when presented at intense illuminance

including wavelengths from the relevant part of the spectrum [62–64]. The differences in

illuminance we measured may correlate with spectral differences; under shaded conditions, the

available light may be reflected from the vegetation or from outside the path of direct sunlight,

and consequently lack red wavelengths (but be rich in greenish or bluish light) [16,17]. Indeed,

previous work modelling *Heliconius* vision suggests that red patterns are more conspicuous to

Heliconius when presented in bright sunlight [41] (though whether this affects *Heliconius*

behaviours has not previously been tested).

Regardless of the proximate mechanism, our data indicate that prevailing light conditions

can influence visual mating behaviours in *Heliconius*. Unlike many other examples, often

involving anthropogenic induced changes in aquatic environments [10–12], the effects we see

here potentially act across very short time and spatial scales. This reflects the forest environment

where the light environment can vary rapidly across both time and space. As such, male response

may depend less on the broader environment (perhaps influenced by forest type), than on the

movement of females across smaller spatial scales (for example between dimly and brightly lit

patches). *Heliconius* butterflies are known to bask in the sun with their wings open, particularly

during the morning hours [42]. This may primarily be for thermoregulation, though other species

are known to display in environments where they are most conspicuous, and *Heliconius* might

follow a similar strategy. *Anolis* lizards, for example, occupy micro-habitats in which their

dewlap colour is most conspicuous [65]. Visual modelling indicates *Heliconius* red patterns are

more conspicuous to avian predators when presented in bright sunlight, suggesting it may also

enhance aposematism [41]. Whatever the ultimate reason for aggregating in sun-exposed patches,

our data suggest that these behaviours could enhance the strength of divergent mating

preferences.

Whether the differences in visual attraction behaviours that we observe translate to

assortative mating, and therefore contribute to reproductive isolation, is an important question.

Previous studies testing for assortative mating between *Heliconius* species in tetrad experiments

similar to ours report low frequencies of interspecific mating [29,66,67]. This was not the case in

our experiments. Nevertheless, the large number of trials in our study allowed us to detect

positive assortative mating, albeit at low levels, though this was much stronger for the 60 trials in

which *H. heurippa* males mated (Fig. 3). Unfortunately, we do not have illuminance data for

these experiments, and this would be difficult to measure given the movement of females.

Considering that the live females may have been basking in the sun, that overall activity of

Heliconius is highest when it is sunny [68] and that mating in *Heliconius* seems to occur more

frequently on sunny than on cloudy days (M. Linares, pers. obs.), it is likely that a large

proportion of the trials with mating outcomes included sun-exposed females. When accounting

for the more frequent involvement of *H. heurippa* males in mating, the mating rates between *H.*

heurippa and *H. t. linaresi* in our tetrad experiments closely match estimates derived from our

experiments with dead mounted females in bright light. Although we cannot rule out a role for

cues acting across other sensory modalities, differences in visual attraction alone are sufficient to

explain the levels of assortative mating we observe.

Considering our data for hybrids, it is overall difficult to ascertain patterns of dominance

for visual attraction. This is perhaps not surprising given that the differences in behaviour

between populations are subtle and are shaped by considerable variation. However, segregation

of the *optix* alleles, which control red pattern elements in *Heliconius* [46], in the backcross to *H.*

t. linaresi did allow us to test for linkage between the warning pattern cue and the corresponding

behaviour. Once again, we observed illuminance-induced shifts in visual attraction, so that the

two types of backcross to *H. t. linaresi* males (*i.e.* red/*Bb* vs. non-red/*bb*) differed in behaviour,

but only under higher illuminance conditions. These results are consistent with physical linkage

between behavioural loci and *optix*, as has been shown elsewhere – but with much greater effects

– between the closely related species *H. cydno* and *H. melpomene* [27]. Physical linkage will help

to maintain key genetic associations (*i.e.* LD) between loci underlying ecologically relevant traits

and those for premating isolation, like mating preferences, facilitating speciation with gene flow

in general [21,23], and hybrid trait speciation [69] in particular. Although the differences between

backcross to *H. t. linaresi* genotypes account for ~50% of that observed in the two parental taxa,

in absolute terms, the effects are unlikely to permit sufficient power for a formal QTL study and

caution should be exercised when interpreting our results. Our results are consistent with a simple

genetic mechanism by which behavioural alleles were acquired alongside red colour pattern

alleles through introgression from *H. melpomene* into the '*heurippa/timareta/cydno*' lineage

[69,70]. However, it seems unlikely that the behavioural differences we observe here would

rapidly lead to strong reproductive isolation, as predicted under a model of hybrid trait speciation

[66,69].

In conclusion, the behavioural differences we observe for *H. heurippa* and *H. t. linaresi*

are similar in strength to those reported elsewhere for other *Heliconius* taxa at the early stages of

divergence (e.g. [30,53]). Alone, these may represent only weak barriers to gene flow; however,

by augmenting divergent ecological selection acting on the warning pattern cue, they may

facilitate the accumulation of additional barriers as speciation proceeds. In addition, our results

suggest that the degree to which *Heliconius* warning pattern contribute to premating isolation

may depend on local illuminance, which can change rapidly in both time and space. Without

more stable differences in the sensory environment, perhaps facilitated by shifts in habitat use,

these fluctuations may constrain speciation. However, *Heliconius* are known to aggregate and

display their warning patterns in sun-exposed patches within the broader forest environment, and

our results suggest that this context would enhance premating isolation. Traits predominantly

shaped by ecology frequently act as mating cues, which by coupling divergent natural selection to

premating isolation can promote speciation with gene flow [6]; our results highlight that this

effect may depend on the sensory conditions during which these ecological mating cues are

displayed to conspecifics.

ACKNOWLEDGEMENTS

We are very grateful to the Abondano-Almeida family for helping AEH and MF settle in

Colombia; Juan Sebastián Sánchez, Óscar Penagos, Isabel Leon and Lina Gabriela Melo for

assistance in the insectaries; Annika Neuhaus, Morgan Oberweiser, Saoirse McMahon and Lucas

Asis for help scoring videos; and Martin Küblbeck, Lucie Queste, Matteo Rossi, Daniel Shane

Wright and Stephen Montgomery for comments on the manuscript. Field collections and

insectary rearing were conducted under permit no. 530 issued by the Autoridad Nacional de

Licencias Ambientales of Colombia (ANLA). CS and CP were funded by Universidad del

Rosario (Grant Number: IV-FGD005). ML was partially funded by the Faculty of Natural

Sciences at Universidad del Rosario. AEH, MF and RMM were supported by an Emmy Noether

fellowship and research grant awarded to RMM by the Deutsche Forschungsgemeischaft (DFG)

(Grant Number: GZ: ME 4845/1-1).

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Supplementary material: Light environment influences mating behaviours during the early stages of divergence in tropical butterflies

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1) Supplementary Figures

Figure S1: Examples from processed videos showing 'poorly lit', median and 'brightly lit' *H. heurippa* 'female' (left) and a *H. t. linaresi* 'female' as a comparison at same illuminance (right). Median illuminance at the *H. heurippa* 'female' was calculated from illuminance data at the *H. heurippa* 'female' when 'pure' males showed a response to either model. This median value was used as threshold to build the categorical model for 'pure' males. The exact same procedure was used for BL hybrid males. The specific *lux* values for 'poorly lit' and 'brightly lit' conditions (6'200 and 132'268 lux) shown here were randomly chosen from the bottom and the top quartile of the distribution. The presented frames were picked from different recording days. Additionally to the mounted female, each frame shows a conspecific male approaching/courting the 'female' and dots used to identify individual males are visible on their wings. A change in hue of the red patch on *H. heurippa* seems to be visible under different illuminance conditions, but it should not be forgotten that 1) human vision is vastly different from butterfly vision and 2) that this is a representation produced by the GoPro camera (GoPro, San Matteo, CA), which is post-processing colour-composition of frames.

Figure S2: All interaction terms involving male type from fully saturated models (*i.e.* Table S2, model #3 for *H.*
heurippa and *H. t. linaresi* males and Table S4, model #1 for backcross to *H. t. linaresi* males). **A+B**: Two-way
interactions between illuminance at the *H. heurippa* mounted female and male type. Proportion of interactions with
the *H. heurippa* female of *H. heurippa* (red) and *H. t. linaresi* (blue) males (A; resembles Fig. 2A, but comes from a
different underlying model); and differently coloured backcross to *H. t. linaresi* males (B, red males (red), males
without red (blue)). **C+D**: The same for the two-way interactions between illuminance at the *H. t. linaresi* mounted
female and male type **E-H**: Three-way interactions between male type and the two illuminance measures. Male type
is indicated on top of each graph. Colouration of each region in the graph depends on predicted preference for the *H.*
heurippa mounted female at the given light conditions (scale depicted on right). White contour lines show Kernel
densities weighted by preference score, which essentially gives the same information as the colour gradient ('peaks'
are where the landscape is most red). All illuminance-axes are logarithmically scaled. Dashed lines in G and H show
median illuminance measures at the respective axis, used to categorize conditions into 'poorly' and 'brightly' lit for
the categorical models.

Figure S3: **A-D**: Raw data points of *H. t. linaresi*, *H. heurippa*, non-red and red backcross to *H. t. linaresi* males (from top to bottom) in a two-dimensional illuminance space. Horizontal dimension shows illuminance at the mounted *H. heurippa* female, vertical dimension shows illuminance at the mounted *H. t. linaresi* female. Blue dots indicate responses of males to the mounted *H. t. linaresi* female and red dots indicate responses to the mounted *H. heurippa* female. Kernel densities are shown for each set of data (blue lines for responses to *H. t. linaresi*, red lines for responses to *H. heurippa*). **E-H**: the two-dimensional illuminance space was then divided into 15 squares for each male type and local preferences within each square were calculated. Graphs directly relate to graphs A-D. The higher the preference for the mounted *H. heurippa* female in a square, the more red the square; the higher the preference for the *H. t. linaresi* female, the more blue (see scale on the right). Squares without response are white. These local preferences have to be interpreted in combination with the local sample sizes (as shown in left column), since they often rely on very few observations. All axes are logarithmically scaled.

Figure S4: Individual trajectories from categorical models. Visual preferences of pure males and hybrid males are
displayed by type. Within each type, data is split depending on illuminance around the *H. heurippa* mounted female,
into poorly and brightly lit. Whenever an individual appeared under both light conditions, it is connected by a
reaction norm line in this plot. Transparency of reaction norm lines within each panel is scaled by the maximum
number of observations available for one individual within this panel. For *H. t. linaresi*, *H. heurippa* and BL males,
posterior means for each EMM are calculated from the same underlying categorical models as used for Fig. 2 and
Fig. 4, respectively. For F_1 and BH males, a categorical model was fit including male type, the illuminance category
around the *H. heurippa* 'female' (determined again by the median measurement) and their interaction. EMMs were
then extracted using the same logic as for the other types. The difference between two groups (in this case between
poorly and brightly lit conditions within each type) is shown by the same Gardner-Altman plot type as used for Fig. 2
and Fig. 4. Small red and blue line located at the right side of each panel show estimators for *H. t. linaresi* and *H.*
heurippa males from Fig. 2B as a reference. [*Figure follows on next page*]

Figure S5: Relative attraction towards *H. heurippa* 'females' (i.e. the proportion of interactions with the mounted *H.*
heurippa 'female' as opposed to the *H. t. linaresi* 'female') for non-red (*bb*) BL males (blue dots) and red (*Bb*) BL
males (red dots). Data and estimators are across all light environments. Note that only preference data with
associated illuminance data were included in the underlying statistical models (a few of the raw data points displayed
here therefore were not considered by the underlying model). The difference between the two groups is shown with
the same logic as used for Fig. 2 and Fig. 4. Small red and blue line located at the right side of each panel show
estimators for *H. t. linaresi* and *H. heurippa* males from Fig. 2B as a reference.

Table S1: Collection locations of *H. t. linaresi* and *H. heurippa*.

Taxon	Location
Heliconius timareta linaresi	Guayabal (2°41'04"N, 74°53'17"W)
Heliconius heurippa	Lejanías (03°34'0"N, 74°04'20.4"W)
	Buenavista (4°10'30"N, 73°40'41"W)
	Santa María (04°53'28.2"N, 73°15'11.4"W)

Table S2: Information on different experimental hybrid lines. Left column shows F_1 broods from which lines originated. For the third line, we summarized two F_1 crosses and their subsequent crosses into one hybrid line. These two F_1 crosses had the same *H. heurippa* father, but a different *H. t. linaresi* mother. Individual counts in brackets show how many males of each brood were tested in the experiment. 32 other F_1 males from 7 broods were tested in the experiment and are not mentioned here, as their brood was not involved in generating backcrosses (13 individuals/5 broods of those being from a cross between *H. t. linaresi* mother and *H. heurippa* father and 19 individuals/2 broods being from the reciprocal F_1 cross with *H. heurippa* mother and *H. t. linaresi* father). *H. heurippa* is abbreviated as *heu* and *H. t. linaresi* as *lin* in the table.

F_1 brood	Backcross to *lin* with male F_1	Backcross to *lin* with female F_1	Backcross to *heu* with male F_1
mother: *lin*, father: *heu*	mother: *lin*, father: F_1	mother: F_1, father: *lin*	mother: *heu*, father: F_1
C18_002 (12 indiv.)	7 broods (45 indiv.)	2 broods (9 indiv.)	2 broods (20 indiv.)
C18_020 (6 indiv.)	3 broods (83 indiv.)	1 brood (15 indiv.)	1 brood (9 indiv.)
C18_034 & C18_036 (half-sib.) (13 indiv.)	4 broods (18 indiv.)	2 broods (10 indiv.)	3 broods (6 indiv.)
C19_014 (-)	1 brood (8 indiv.)	-	-

755　**Table S3**: *GoPro Hero 5 Black™* (GoPro, San Matteo, CA) camera settings and equipment. All auto-settings (except

756　were indicated) were disabled.

Settings	
Resolution	1920 x 1080 pixels
Frame rate	60 fps
Shutter speed	1/480
Field of view	*Narrow*
Colour	*GoPro*
Sharpness	*High*
ISO	400
White balance	*Auto*
Equipment	• 128 GB *SanDisk™ Extreme* SD card • *Neewer™ UV* filter (reduced overexposure of yellow bands) • *Xlayer™ 210033* powerbank (charging while recording)

Table S4: 19 possible fixed-effect structures for the pure males of *H.* heurippa and *H. t.* linaresi (fixed effect 'type').
'I_H' and 'I_L' stand for log-illuminance measures at the *H.* heurippa and the *H. t.* linaresi mounted females,
respectively. A '*' sign in a model formula indicates that all involved variables as well as all possible interactions
between them are included. All models include trial and male ID as random effects. Including trial (and in many
cases also individual ID) as a random effect has often been disregarded in previous analyses. However, we found
strong correlation among individuals' behaviour during trials (see supplementary R Markdown), likely due to the
unique combination of males, mounted females and their position in the experimental cage during a given trial. The
latter may have also affected sensitivity of our motion-detection software. We therefore considered it inevitable to
correct for this. For each model we calculated a LOOIC and WAIC value, which mostly agree. Based on the WAIC
value, differences between best model and other models are calculated (ΔWAIC) as well as a model-weight score.

#	Fixed Effects Term	LOOIC	WAIC	ΔWAIC	Weight$_{WAIC}$
1	~ type * I_H	3773.40	3768.52	0.00	0.29
2	~ type * I_H + I_L	3774.63	3769.68	1.16	0.16
3	~ type * I_H * I_L	3775.20	3770.08	1.56	0.13
4	~ type * I_H + I_H * I_L	3775.99	3771.04	2.52	0.08
5	~ type * I_H + type * I_L	3776.53	3771.52	3.01	0.06
6	~ type * I_H + type * I_L + I_H * I_L	3776.62	3771.56	3.04	0.06
7	~ type	3777.82	3773.07	4.55	0.03
8	~ type + I_L	3778.10	3773.20	4.68	0.03
9	~ type + I_H	3778.11	3773.33	4.81	0.03
10	~ I_L	3778.50	3773.62	5.10	0.02
11	~ type * I_L	3778.83	3773.95	5.43	0.02
12	~ 1	3779.24	3774.44	5.92	0.02
13	~ I_H	3779.45	3774.58	6.06	0.01
14	~ type + I_H + I_L	3779.75	3774.87	6.35	0.01
15	~ I_H + I_L	3780.58	3775.58	7.06	0.01
16	~ type + I_H * I_L	3780.77	3775.80	7.28	0.01
17	~ type * I_L + I_H	3780.82	3775.99	7.47	0.01
18	~ type * I_L + I_H * I_L	3781.61	3776.60	8.09	0.01
19	~ I_H * I_L	3781.85	3776.78	8.26	0.00

Male type	Light environment	Mean	Lower	Upper
H. t. linaresi (left in Fig. 2B)	Across all	0.447	0.383	0.513
H. heurippa (right in Fig. 2B)	Across all	0.515	0.449	0.582
H. t. linaresi (left in Fig. 2C)	*H. heurippa* -	0.464	0.391	0.540
H. heurippa (right in Fig. 2C)	*H. heurippa* -	0.456	0.385	0.527
H. t. linaresi (left in Fig. 2D)	*H. heurippa* +	0.437	0.362	0.514
H. heurippa (right in Fig. 2D)	*H. heurippa* +	0.572	0.493	0.650
BL without red (left in Fig. S5)	Across all	0.461	0.414	0.508
BL with red (right in Fig. S5)	Across all	0.493	0.444	0.543
BL without red (left in Fig. 4A)	*H. heurippa* - / *H. t. linaresi* -	0.460	0.406	0.515
BL with red (right in Fig. 4A)	*H. heurippa* - / *H. t. linaresi* -	0.475	0.419	0.531
BL without red (left in Fig. 4B)	*H. heurippa* - / *H. t. linaresi* +	0.477	0.401	0.554
BL with red (right in Fig. 4B)	*H. heurippa* - / *H. t. linaresi* +	0.443	0.359	0.528
BL without red (left in Fig. 4C)	*H. heurippa* + / *H. t. linaresi* -	0.426	0.350	0.505
BL with red (right in Fig. 4C)	*H. heurippa* + / *H. t. linaresi* -	0.538	0.461	0.616
BL without red (left in Fig. 4D)	*H. heurippa* + / *H. t. linaresi* +	0.438	0.384	0.491
BL with red (right in Fig. 4D)	*H. heurippa* + / *H. t. linaresi* +	0.504	0.446	0.563

Estimates based on	*H. heurippa* ♀ *H. t. linaresi* ♂	*H. t. linaresi* ♀ *H. t. linaresi* ♂	*H. heurippa* ♀ *H. heurippa* ♂	*H. t. linaresi* ♀ *H. heurippa* ♂
tetrad experiment	0.168 [0.099; 0.253]	0.157 [0.09; 0.239]	0.405 [0.305; 0.508]	0.27 [0.183; 0.366]
mounted females: across all	0.149 [0.084; 0.23]	0.184 [0.111; 0.27]	0.349 [0.255; 0.45]	0.329 [0.236; 0.429]
mounted females: poorly lit *H. heurippa*	0.152 [0.086; 0.233]	0.181 [0.109; 0.266]	0.315 [0.224; 0.414]	0.363 [0.267; 0.465]
mounted females: brightly lit *H. heurippa*	0.146 [0.082; 0.226]	0.187 [0.114; 0.273]	0.387 [0.289; 0.489]	0.291 [0.203; 0.389]

Table S7: 19 possible fixed-effect structures for the BL males (fixed effect 'type' refers here to wing colour
phenotype of hybrids). 'I_H' and 'I_L' stand for log-illuminance measures at the *H. heurippa* and the *H. t. linaresi*
mounted females, respectively. A '*' sign in a model formula indicates that all involved variables as well as all
possible interactions between them are included. All models include trial, male ID and brood as random effects. Each
model structure has a LOOIC and WAIC value, which mostly agree. Based on the WAIC value, differences between
best model and other models are calculated (ΔWAIC) as well as a model-weight score.

#	Fixed Effects Term	LOOIC	WAIC	ΔWAIC	Weight$_{WAIC}$
1	~ type * I_H * I_L	6815.06	6811.69	0.00	0.38
2	~ type * I_H + type * I_L + I_H * I_L	6815.98	6812.60	0.91	0.24
3	~ type * I_H + type * I_L	6817.15	6813.84	2.15	0.13
4	~ type * I_H + I_H * I_L	6817.82	6814.43	2.74	0.10
5	~ type * I_H	6820.15	6816.86	5.17	0.03
6	~ type * I_H + I_L	6820.78	6817.48	5.79	0.02
7	~ type + I_H * I_L	6821.33	6817.96	6.27	0.02
8	~ I_H * I_L	6822.03	6818.67	6.98	0.01
9	~ type	6821.94	6818.68	6.99	0.01
10	~ type * I_L + I_H * I_L	6822.28	6818.95	7.27	0.01
11	~ 1	6822.46	6819.18	7.50	0.01
12	~ type + I_H	6822.96	6819.70	8.01	0.01
13	~ type + I_L	6823.18	6819.94	8.26	0.01
14	~ I_H	6823.50	6820.19	8.50	0.01
15	~ I_L	6823.80	6820.52	8.84	0.00
16	~ type + I_H + I_L	6824.36	6821.03	9.35	0.00
17	~ type * I_L	6825.21	6821.92	10.23	0.00
18	~ I_H + I_L	6825.20	6821.93	10.24	0.00
19	~ type * I_L + I_H	6825.55	6822.23	10.54	0.00