



Continuous Variation in an Aposematic Pattern Affects Background Contrast, but Is Not Associated With Differences in Microhabitat Use

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Variation in aposematic signals was once predicted to be rare, yet in recent years it has become increasingly well documented. Despite increases in the frequency with which polytypism and polymorphism have been suggested to occur, population-wide variance is rarely quantified. We comprehensively sampled a subpopulation of the poison frog *Oophaga sylvatica*, a species which is polytypic across its distribution and also shows considerable within-population polymorphism. On one hand, color pattern polymorphism could be the result of multifarious selection acting to balance different signaling functions and leading to the evolution of discrete sub-morphs which occupy different fitness peaks. Alternatively, variance could simply be due to relaxed selection, where variation would be predicted to be continuous. We used visual modeling of conspecific and heterospecific observers to quantify the extent of within population phenotypic variation and assess whether this variation produced distinct signals. We found that, despite considerable color pattern variation, variance could not be partitioned into distinct groups, but rather all viewers would be likely to perceive variation as continuous. Similarly, we found no evidence that frog color pattern contrast was either enhanced or diminished in the frogs' chosen microhabitats compared to alternative patches in which conspecifics were observed. Within population phenotypic variance therefore does not seem to be indicative of strong selection toward multiple signaling strategies, but rather pattern divergence has likely arisen due to weak purifying selection, or neutral processes, on a signal that is highly salient to both conspecifics and predators.

Keywords: aposematism, crypsis, poison frog, *Oophaga sylvatica*, visual modeling

INTRODUCTION

Aposematic (warning) signals evolve to convey important information related to prey defenses to potential predators (Stevens and Ruxton, 2012; Caro and Ruxton, 2019). Predators evolve, or learn, to associate certain prey characteristics with chemical or physical defenses and subsequently avoid prey bearing these signals (Stevens and Ruxton, 2012; Caro and Ruxton, 2019). By the

simplest functional interpretation of aposematism, warning signals should be readily recognizable, reliable, and memorable to facilitate rapid associative learning (Mappes et al., 2005; Stevens and Ruxton, 2012). Positive frequency dependent selection is, therefore, expected to drive further homogenization of signal characteristics with common local phenotypes being avoided more consistently than rare phenotypes (Mappes et al., 2005; Chouteau and Angers, 2011; Chouteau et al., 2016).

In nature, however, aposematic signals can vary considerably between individuals of a single species, both within and between populations (Briolat et al., 2019). Considerable attention has been paid to how warning coloration evolves and functions (Caro and Ruxton, 2019), yet basic questions related to polymorphism and polytypism remain. Does variation serve an adaptive function such as local specialization to distinct environmental conditions, or is it merely permitted due to a relaxation in selection where neutral processes can lead to signal divergence? Once variation has arisen in allopatry, it may then be reinforced through mechanisms related to local predation pressure, assortative mating or intraspecific agonistic interactions, or a combination of several factors, which can further drive divergence between genetically and physically isolated populations (Gray and McKinnon, 2007; Briolat et al., 2019). Variation can, however, also arise in sympatric populations without physical barriers to gene flow (Rojas, 2017; Briolat et al., 2019). The mechanisms underlying the evolution and persistence of this variance are, however, less well understood, and in many cases it is still unclear whether such variation has evolved for an adaptive function.

The characteristics of animal color patterns are shaped by the simultaneous influence of multiple selection pressures, leading to potential trade-offs rather than the optimization of single functions (Endler and Mappes, 2004; Stevens, 2007; Stevens and Ruxton, 2012; Briolat et al., 2019). For example, individual components of aposematic signals can be locally adapted for saliency when viewed against particular microhabitats, serve to incorporate additional defensive signals, or contain traits used in sexual selection. Based on its primary function, each element could theoretically have its own distinct optimal form which influences fitness under specific sources of selection. Polymorphism can evolve due to adaptive benefits, whereby different forms arise as a by-product of unique adaptive processes (Briolat et al., 2019). Alternatively, instead of being fostered by selection, polymorphism may simply be permitted where purifying selection is weak, or optimal signal efficacy is difficult to achieve, leading to variation arising *via* neutral processes (Briolat et al., 2019). With such distinct plausible mechanisms at play, it is perhaps not unsurprising we have a weak overall understanding of why intra-population variance in aposematic signals exists.

Some of the most salient examples of phenotypic variation are found in the Neotropical poison frogs (Dendrobatidae), which exhibit notable examples of both discrete and continuous variation in color and pattern, found both in sympatry (polymorphism) and allopatry (polytypism) (Summers et al., 2003; Maan and Cummings, 2012; Rojas, 2017). Color patterns have evolved both for defensive and communicative functions, and in many cases under the influence of multiple

selection pressures (Summers et al., 1999; Saporito et al., 2007; Maan and Cummings, 2008, 2009, 2012; Wollenberg et al., 2008; Tazzyman and Iwasa, 2010; Crothers et al., 2011; Cummings and Crothers, 2013; Rudh, 2013; Crothers and Cummings, 2015; Dreher et al., 2015). Consequently, signals with multiple components or functions have arisen due to context specific interactions between sources of selection and/or neutral processes (Wollenberg et al., 2008; Tazzyman and Iwasa, 2010; Crothers et al., 2011; Crothers and Cummings, 2013, 2015; Cummings and Crothers, 2013; Barnett et al., 2018). For example, aposematic signals have been co-opted and exaggerated by intraspecific communication (Maan and Cummings, 2008, 2009), can blend together when viewed from a distance to act as camouflage (Barnett et al., 2018), and can disrupt an observer's ability to track escape movements (Hämäläinen et al., 2015).

In poison frogs, discrete color forms have evolved in geographically isolated populations as founder effects and honest signaling of local resource availability have been reinforced through sexual imprinting, selective mate choice, and predator learning (Summers et al., 1999, 2003; Darst et al., 2006; Reynolds and Fitzpatrick, 2007; Maan and Cummings, 2008, 2009, 2012; Wang and Shaffer, 2008; Tazzyman and Iwasa, 2010; Crothers et al., 2011; Richards-Zawacki and Cummings, 2011; Crothers and Cummings, 2013; Hegna et al., 2013; Yang et al., 2019). For example, in *Oophaga pumilio*, where dietary derived toxicity is low, selection for camouflage has outweighed selection for conspicuous signaling, and more cryptic colors and behaviors have evolved (Maan and Cummings, 2012; Rudh et al., 2012; Rudh, 2013). Within color morphs of *O. pumilio*, continuous variation in color saturation and brightness have been associated with intraspecific communication, with brighter individuals more likely to win physical contests and attract mates (Crothers et al., 2011; Crothers and Cummings, 2013, 2015; Dreher et al., 2017). Moreover, in *O. pumilio* certain phenotypes and behaviors are correlated with microhabitats which enhance signal efficacy (Pröhl and Ostrowski, 2011; Willink et al., 2014), although they do not appear to be behaviorally manipulating saliency on fine scales (Dugas et al., 2020). In another polymorphic and polytypic poison frog species, *Dendrobates tinctorius*, sympatric color variation can also correlate with specialization in specific microhabitats, or behavioral syndromes that optimize signal saliency within their chosen microhabitats (Rojas et al., 2014a,b; Rojas, 2017).

The poison frog *Oophaga sylvatica* shows polytypic variation across its range as well as remarkable within-population polymorphism (Roland et al., 2017). Limited insights into color pattern evolution can be gleaned from population-level genetic assessments, which indicate well-defined clades broadly split between north and south, but also reveal large geographic regions with high levels of genetic admixture, including within polymorphic populations (Roland et al., 2017). Polymorphic populations are proposed to either be the result of admixture between phenotypically distinct adjacent populations in secondary contact, or due to a range extension from a polymorphic population into adjacent populations (Roland et al., 2017). Either scenario suggests that color patterns may not be under strong selection within the northern clade populations,

and/or that coloration does not play a significant role in the geographic structure of phenotypically divergent *O. sylvatica* populations across at least substantial portions of its distribution.

We explored the role of phenotypic variation in a highly variable population of *O. sylvatica* found in northwestern Ecuador. These frogs display high contrast dorsal patterns that are highly distinct from their leaf litter substrates (Yeager and Barnett, 2020). However, as early as in the formal description of *O. sylvatica* it has been noted that human observers could easily confuse frogs from a similar red/black morph with the forest floor (Funkhouser, 1956), suggesting a potential balance between camouflage and conspicuous signaling.

We used computational visual models, representing the vision of diverse predator classes and conspecifics, to study the perception of phenotypic variation by sources of natural and sexual selection. We first addressed whether frog variation was continuous, or if it could be grouped into visually distinct clusters, and then quantified which components of the phenotype (chromatic (hue), achromatic (brightness), or pattern contrast) best defined phenotypic variation. Next, we investigated whether variance between the aposematic signals of individual frogs correlated with the microhabitats in which they were found to either maximize or minimize visual contrast. We expected that regardless of whether variation is continuous or discrete, if it has evolved to facilitate distinct behavioral, defensive, or reproductive strategies then it should be associated with differences in saliency based on microhabitat use (Rojas et al., 2014a,b; Rojas, 2017; Briolat et al., 2019; Barnett et al., 2021a). Conversely, variation may be permitted under relaxed selection, and attributable to non-adaptive processes such as drift. In such instances we would expect variation to be continuous and not associated with signaling differences within their occupied microhabitats. Deciphering how this variation arises and is maintained within a single population has important implications for our understanding of how predators generalize behaviors across variable aposematic signals, how multiple selection pressures interact, how the evolution of discrete polymorphism may arise from a monomorphic ancestor, and the evolutionary implications of polymorphism (Gray and McKinnon, 2007; Briolat et al., 2019).

MATERIALS AND METHODS

Data Collection

In March 2020, we photographed 35 *Oophaga sylvatica* (Perla morph) at the private forest reserve “Bosque Protector la Perla” near La Concordia, Ecuador. This represented a comprehensive survey of every individual observed in the area by a team of five experienced observers, over 2 days. The Perla morph is approximately 26 mm in length and is predominantly black with a red pattern that varies from small spots to larger irregular blotches, and to whole patches of homogenous color (Figure 1). Previous work suggests that the frogs’ colors have very high internal and external contrast, and that UV

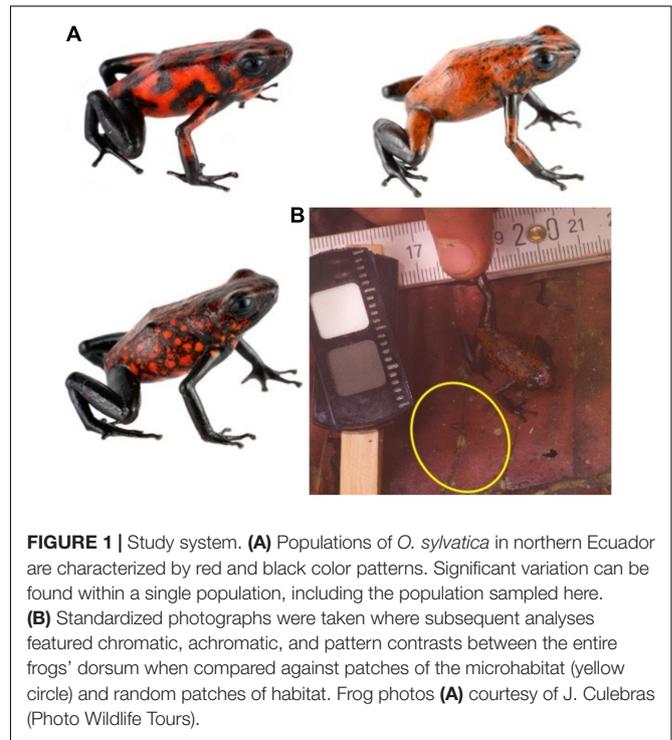


FIGURE 1 | Study system. **(A)** Populations of *O. sylvatica* in northern Ecuador are characterized by red and black color patterns. Significant variation can be found within a single population, including the population sampled here. **(B)** Standardized photographs were taken where subsequent analyses featured chromatic, achromatic, and pattern contrasts between the entire frogs’ dorsum when compared against patches of the microhabitat (yellow circle) and random patches of habitat. Frog photos **(A)** courtesy of J. Culebras (Photo Wildlife Tours).

reflectance is minimal both from the frogs and their natural leaf litter background (Yeager and Barnett, 2020). Frogs were spotted using exhaustive visual surveys. We cannot ensure that frog orientation was not altered due to the presence of the observers, however, the background is isotropic, and we used the location where the frog was first spotted as the microhabitat context in which potential predators would process frogs’ visual signals.

We photographed each frog within the microhabitat where it was first observed (Figure 1B), following methods detailed in Yeager and Barnett (2020, 2021). In brief, each image was taken using a quartz converted UV sensitive Canon EOS 7D camera combined with a metal body NIKKOR EL 80 mm lens. As UV reflectance is negligible in this population ($n = 4$ tested in Yeager and Barnett, 2020) we did not include UV wavelengths and only utilized images in the human visible range (400 – 700 nm). The camera was mounted on a tripod and the lens was fitted with a Baader UV-IR blocking filter that allowed transmission from 420 to 680 nm. All photographs were taken under natural ambient lighting, each image included a 10% and a 77% neutral reflectance standard, and we saved all photographs in RAW format.

Image Processing and Visual Modeling

We used the MICA toolbox (Troscianko and Stevens, 2015) to standardize and scale the images in ImageJ v1.52k (Schneider et al., 2012). From each image, we selected regions of interest (ROIs) from within the red and black patches of the frog, around the frog’s whole body, and around a patch of adjacent substrate of a size equal to, or slightly greater than, that of the frog (Figure 1). We then assessed color and

pattern using three well characterized computational models of visual perception representing important sources of natural selection (predatory birds and snakes) and sexual selection (conspecific frogs).

For our model of bird vision, we used the tetrachromatic visual system of the Eurasian blue tit (*Cyanistes caeruleus*, Paridae). The blue tit has four single cones (λ_{max} : UV = 372 nm, SWS = 413 nm, MWS = 508 nm, and LWS = 573 nm) that populate the retina with a ratio of 1:2:2:3, and one double cone (λ_{max} : D = 565 nm) (Hart et al., 2000; Hart, 2001). As UV was negligible, we excluded these wavelength (<400 nm) to produce a tetrachromatic VIS-sensitive model which covered 400–700 nm (Yeager and Barnett, 2020, 2021). The snake model used the trichromatic vision of the coachwhip (*Masticophis flagellum*, Colubridae) and included three single cones (λ_{max} : UV = 362 nm, MWS = 458 nm, and LWS = 561 nm) with a ratio of 17:2:1 (Macedonia et al., 2009; Maan and Cummings, 2012). To represent the frogs' own visual system, we used the trichromatic vision of the closely related *O. pumilio* (Dendrobatidae). The *O. pumilio* visual model included three single cones (λ_{max} : SWS = 466 nm, MWS = 489 nm, and LWS = 561 nm) with a cone ratio of 4:3:1 (Siddiqi et al., 2004; Maan and Cummings, 2012).

We calculated chromatic (ΔS) and achromatic (ΔL) contrast between the red and black regions of each frog (internal contrast), and between the mean of the whole frog and the mean of each background (external contrast), using the log-linear receptor noise limited model implemented through the MICA toolbox (Vorobyev et al., 1998; Vorobyev and Osorio, 1998; Troscianko and Stevens, 2015). Chromatic contrast was calculated using the responses of the single cones and we calculated achromatic contrast from the response of the D cone in the bird model and from the LWS cones in the snake and frog models. As specific predator species which could influence frog coloration are not known for *O. sylvatica*, and to keep our results comparable to our previous studies (Yeager and Barnett, 2020, 2021; Barnett et al., 2021b) all Weber fractions were set at 0.05. The model computes visual discrimination in a manner equivalent to “just noticeable differences” (JNDs) where higher values indicate greater ease of color discrimination. A score of 1 is the absolute discrimination threshold defined by intrinsic noise within the photoreceptor and a score of 3 is a realistic discrimination threshold under natural lighting conditions (Vorobyev and Osorio, 1998).

Pattern energy was calculated over each of the whole frog ROIs and each of the background ROIs using Fast Fourier bandpass filtering (granularity analysis) from the achromatic channel of each visual model in the MICA toolbox. We measured pattern energy as the standard deviation of the pixel values for five filter bands that doubled in spatial frequency at each step from 16 px (0.5 mm) to 256 px (8.0 mm). To compare pattern energy distributions between the frogs and their backgrounds we calculated the area between pattern energy curves using a piecewise linear function (function *approxfun*) from base R v4.0.5 (R Core Team, 2021). Hereafter, the area between frog and background pattern energy curves is referred to as “pattern contrast,” higher

values indicate that the frog's pattern was more distinct from the background.

Quantifying Intra-Population Variation

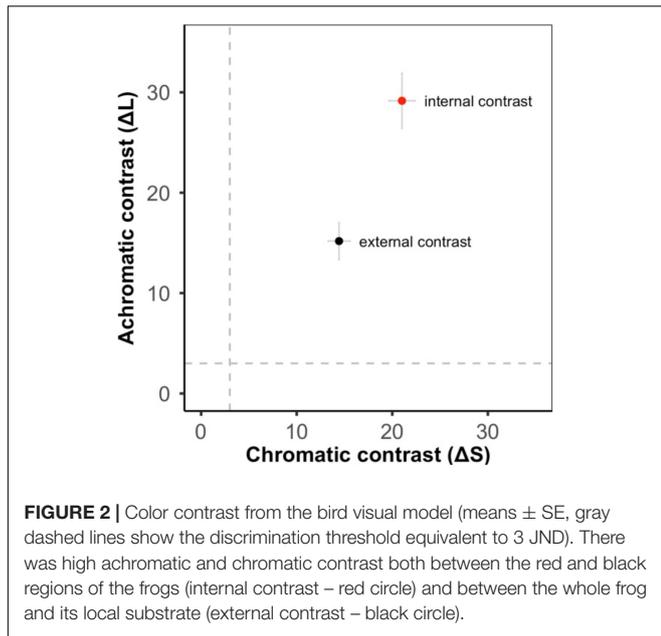
To examine whether frog color pattern variation was continuous or divided into visually distinct sub-morphs we used *k*-means clustering and factor analysis in R v4.0.5 (Barnett et al., 2021a; R Core Team, 2021). For each of the frogs, we used the internal chromatic (ΔS) and achromatic (ΔL) contrast as well as the pattern energy at each of the five spatial scales (0.5, 1.0, 2.0, 4.0, and 8.0 mm). In the *k*-means clustering, we used the gap statistic (function *clusGap*) from R package *cluster* (Maechler et al., 2021), to estimate the optimal number of clusters (Tibshirani et al., 2001). We set the maximum number of clusters to 10 and ran 5000 Monte Carlo bootstrap samples to compute our reference distribution. We then interpreted the smallest local maximum in gap score as the optimal number of clusters (Tibshirani et al., 2001; Maechler et al., 2021). Next, we used factor analysis with two factors and varimax rotation (function *factanal*) from base R v4.0.5 (R Core Team, 2021) to investigate which parameters of the color pattern contributed the most to intra-population variation (Barnett et al., 2021a). We repeated these analyses separately for the bird, snake, and frog visual models. Two frogs were removed from the snake analysis due to image calibration issues.

Background Contrast

We next asked whether frog variation affected contrast against the background. As the majority of the frogs' variation was found in the distribution of pattern (see *Factor 1* in Results: Intra-population variation), and as the red and black are highly distinct from the background (Yeager and Barnett, 2020; **Figure 2** and **Supplementary Figures 1–3**), we focused on how pattern variation affects background pattern matching and distance-dependent color blending by comparing the whole frog ROIs to their local backgrounds. We compared the mean hue, mean luminance, and pattern energy distribution of each frog to that of the microhabitat where it was first encountered (external chromatic, achromatic, and pattern contrast), and then ran a series of linear models comparing each form of external contrast to Factor 1 from our factor analysis (function *lm*) using base R v4.0.5 (R Core Team, 2021). The suitable approximation of model assumptions was checked by plotting the distribution of the model residuals, and the significance of the dependent variables was estimated against a null model by the F statistic. A significant relationship would suggest that frog variation is associated with differences in background matching, whereas a non-significant relationship would suggest that any differences in external contrast were affected by background heterogeneity, independent from frog variation.

Background Choice

To test whether frogs were associated with microhabitats that either minimized or maximized the contrast of their own color and pattern characteristics we compared mean achromatic, mean chromatic, and pattern contrast between each frogs' local microhabitat and the alternate microhabitats where the other 34



individuals were observed. We calculated the median contrast of the 34 alternate backgrounds for each frog and compared local contrast to alternate contrast using a series of paired-sample Wilcoxon tests in base R v4.0.5 (R Core Team, 2021). For each visual model, we ran separate tests for achromatic, chromatic, and pattern contrast. A significant result would suggest that frogs are associated with specific microhabitats.

RESULTS

Intra-Population Variation

Color patterns were overall found to be highly contrasting, both in terms of within-frog contrast (e.g., red versus black

TABLE 1 | Percent variance explained by factor analysis for each visual model.

Variance explained (%)	Bird	Snake	Frog
Factor 1	55.7	56.4	55.8
Factor 2	20.2	31.5	19.6
Cumulative variance	75.8	87.9	75.4

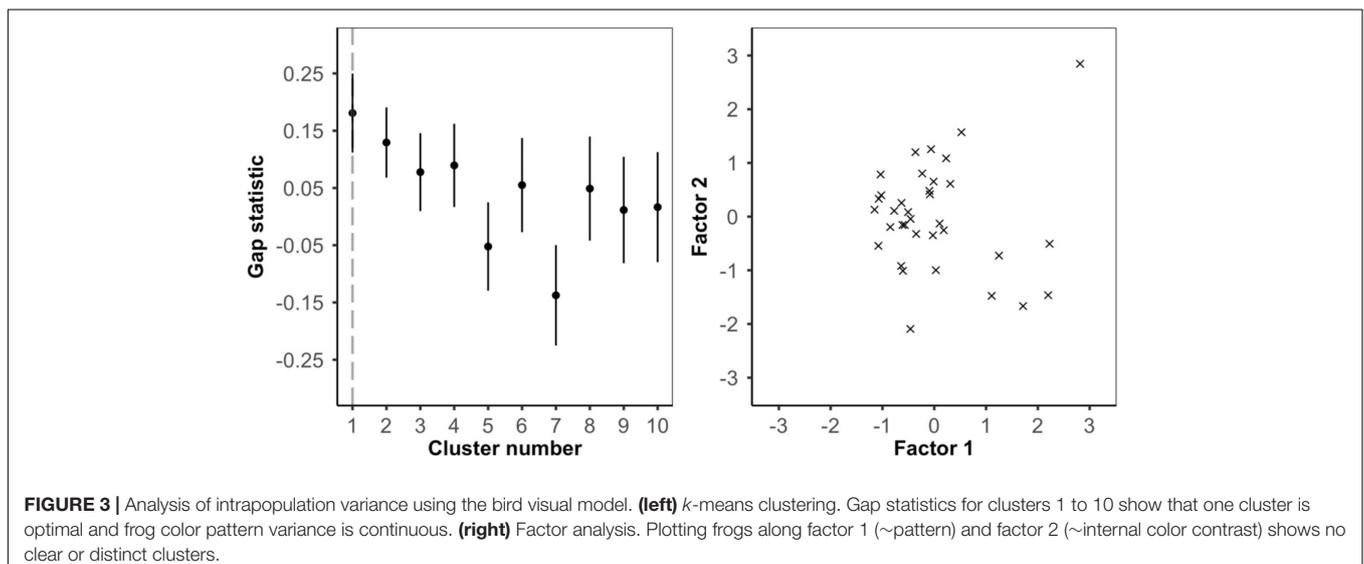
In each visual model the majority of variance is explained by Factor 1 which corresponds to pattern energy across all spatial scales. Bold values significant ($p < 0.05$).

patches) and between the frogs and their microhabitats (Figure 2 and Supplementary Figures 1–3). In the k -means clustering, we found that for each visual model (bird, snake, and frog) the optimal number of clusters was one, suggesting that frog variation was continuous and not divided into discrete sub-morphs (Figure 3 and Supplementary Figures 4, 5). Factor analysis revealed that for each visual model two factors were sufficient to explain $>75\%$ of frog variance (Table 1). Factor 1, which explained $>55\%$ of variance for each model, weighted heavily toward pattern across all size classes. Factor 2, which explained $<32\%$ of the variance in each model was more variable. In the bird and snake models Factor 2 (bird = 20.2% variance, snake = 31.5% variance) mostly corresponded to internal chromatic and achromatic contrast, whereas in the frog model Factor 2 (19.6% variance) predominantly corresponded to low spatial frequency patterning.

Background Contrast

When using the bird and snake visual models, we found that there was no significant relationship between Factor 1 (\sim pattern) and either external achromatic or chromatic contrast. However, there was a significant positive relationship between Factor 1 and external pattern contrast, where frogs with higher Factor 1 scores were more distinct from their local backgrounds (Figure 4; Table 2; Supplementary Figures 6, 7; Supplementary Table 1).

In the frog visual model, we found no significant relationship between Factor 1 and achromatic contrast, but there was a



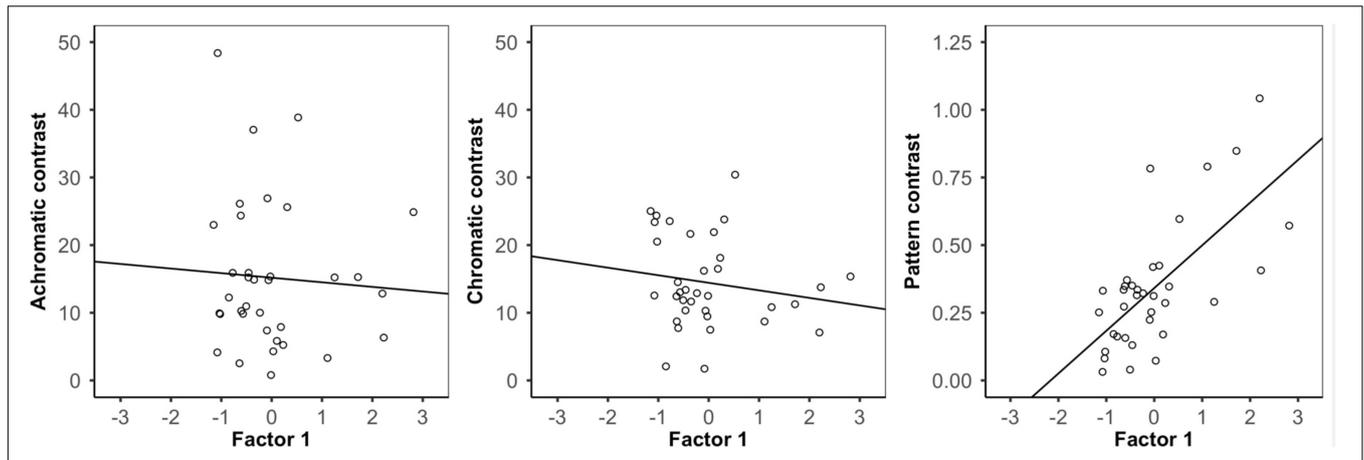


FIGURE 4 | Analysis of background contrast from the bird visual model. Correlation between Factor 1 (pattern variation) score and the achromatic (left), chromatic (middle), and pattern (right) contrast of each frog and its local background. There was no relationship between frog variation and either achromatic or chromatic contrast, but there was a significant relationship between frog pattern and pattern contrast where frogs with higher Factor 1 scores had greater pattern contrast against their local backgrounds.

significant relationship between Factor 1 and both chromatic and pattern contrast. Frogs with higher Factor 1 scores were more distinct from their local backgrounds in both chromatic contrast and patterning, although chromatic contrast remained well above the discrimination threshold in both conditions (Table 2).

Effects of Occupied Microhabitats on Frog Signals

Frogs were most commonly found in leaf litter ($n = 27$) or green vegetation ($n = 8$) substrates. We found no significant effect of background type (local vs. alternative) on frog contrast against the background in the bird and frog visual models (Table 3). Similarly, in the snake model we found no significant effect of background type on achromatic and pattern contrast, but there was a marginally significant effect for chromatic contrast ($V = 170, p = 0.048$; Table 3).

TABLE 2 | Statistical results from the linear models comparing achromatic, chromatic, and pattern contrast against their local microhabitat substrate and Factor 1 from the factor analysis, which explained >55% of total frog variance.

	Bird	Snake	Frog
Achromatic contrast	Adj. $R^2 = -0.026$ $F_{1,33} = 0.13,$ $p = 0.723$	Adj. $R^2 = -0.028$ $F_{1,33} = 0.11,$ $p = 0.737$	Adj. $R^2 = -0.028$ $F_{1,33} = 0.08,$ $p = 0.781$
Chromatic contrast	Adj. $R^2 = -0.001$ $F_{1,33} = 0.94,$ $p = 0.340$	Adj. $R^2 = -0.008$ $F_{1,33} = 0.75,$ $p = 0.392$	Adj. $R^2 = 0.103$ $F_{1,33} = 4.89,$ $p = 0.034$
Pattern contrast	Adj. $R^2 = 0.434$ $F_{1,33} = 27.05,$ $p < 0.001$	Adj. $R^2 = 0.600$ $F_{1,33} = 48.91,$ $p < 0.001$	Adj. $R^2 = 0.447$ $F_{1,33} = 28.48,$ $p < 0.001$

Significant relationships suggest that frog variation affects contrast against the background. Conversely, non-significant relationships suggest that differences in contrast to the background are independent of frog variation and result from background heterogeneity. Bold values significant ($p < 0.05$).

DISCUSSION

The evolution and maintenance of intraspecific variation in aposematic signals continues to draw considerable interest (Rojas, 2017; Briolat et al., 2019). Although in *O. sylvatica* the extent of color pattern variation found within populations is not as striking as that observed between populations (Roland et al., 2017), resolving basic questions related to the degree and organization of phenotypic divergence is an essential step in beginning to understand broader scales of both polymorphism and polytypism in aposematic species (Briolat et al., 2019). A central prediction of within-population polymorphism is that if variance exists, it should be correlated with differences in signal saliency to serve distinct adaptive roles, such as in separate microhabitats within heterogeneous landscapes (Gray and McKinnon, 2007), or where a single color pattern serves divergent anti-predator and social functions (Cummings and Crothers, 2013).

Using a population-wide sampling effort, we sought to understand fine-scale phenotypic variance. We found intra-population color pattern variation in the Perla population

TABLE 3 | Statistical results from the paired-sampled Wilcoxon tests comparing achromatic, chromatic, and pattern contrast between local and alternate background types.

	Bird	Snake	Frog
Achromatic contrast	$V = 268, p = 0.451$	$V = 208, p = 0.201$	$V = 237, p = 0.207$
Chromatic contrast	$V = 233, p = 0.184$	$V = 170, p = 0.048$	$V = 220, p = 0.128$
Pattern contrast	$V = 340, p = 0.692$	$V = 317, p = 0.525$	$V = 337, p = 0.728$

A significant relationship suggests that frogs are associated with backgrounds that either increase or decrease contrast more than the median alternative background. Bold values significant ($p < 0.05$).

of *O. sylvatica* to be continuous, rather than discrete, and largely driven by pattern elements, rather than chromatic or achromatic contrast. Moreover, although individuals differed in pattern distinctness from the background, we found no strong evidence that different individuals were associated with particular microhabitats, such as if they were selecting signaling environments that would either enhance or reduce signal contrast. Taken together, our data suggest that frogs exhibit highly variable patterns which still maintain consistent chromatic and achromatic contrast. However, although seemingly highly variable, the frogs' aposematic signals do appear constrained to a small range of colors such that any variance is likely still perceived as within the same aposematic grouping. Our data are, therefore, consistent with a single fitness peak (or perhaps a broad plateau), with variation arising as a product of weak purifying selection, honest signaling, and/or neutral drift on some elements of the color pattern.

In aposematic signals both color and pattern have been found to convey important information to potential predators (Stevens and Ruxton, 2012; Caro and Ruxton, 2019). Prey bearing patterns comprised of multiple highly contrasting elements have higher survival than prey with single colors (Preißler and Pröhl, 2017). However, in predator education, hue and brightness appear to play a more critical role than the particular arrangement of the pattern (Aronsson and Gamberale-Stille, 2008; Kazemi et al., 2014; Sherratt et al., 2015). In *O. pumilio*, for example, color and pattern are both involved in aversion learning and when controlling for color, larger spotted patterns have been found to be more salient and more likely to be avoided than smaller spotted or plain patterns (Hegna et al., 2011; Qvarnström et al., 2014; Preißler and Pröhl, 2017; Barnett et al., 2021b). However, under natural conditions, both color and pattern can be highly divergent without necessarily increasing predation risk (Hegna et al., 2011; Richards-Zawacki et al., 2013).

The arrangement of pattern elements can also allow seemingly conspicuous signals to incorporate aspects of camouflage, for example, dual signaling as a function of observer viewing distance or by context dependent disruptive coloring (Marshall, 2000; Tullberg et al., 2005; Barnett and Cuthill, 2014; Honma et al., 2015). Distance-dependent signaling takes advantage of limitations in observer visual perception such that when viewed from a distance a high contrast pattern blends together to match the background (Marshall, 2000; Tullberg et al., 2005; Barnett and Cuthill, 2014; Caro et al., 2016). Although the contrast between the average color of the frogs and the background (external contrast) was lower than that found within the frogs' pattern (internal contrast), external chromatic and achromatic contrast was still high. We therefore found no strong evidence that this population of *O. sylvatica* are utilizing distance-dependent pattern blending. Alternatively, in disruptive camouflage, high contrast patterns can interfere with the visual assessment of shapes and edges (Stevens et al., 2006; Stevens and Merilaita, 2009). We did not explicitly test for disruptive coloring but unlike predictions for aposematism, disruptive camouflage can be more effective

when sympatric individuals vary in the expression of pattern characteristics (Troscianko et al., 2021), and future studies could explore this alternative. With this in mind, it is important to note that to test for differences in microhabitat use, we only characterized the local patch of substrate immediately surrounding the frogs. However, we emphasize these were the very patches where frogs were initially detected, representing ecologically relevant contexts. Yet when the whole complexity of the visual environment is considered, these frogs may be more difficult to detect under certain conditions than currently recognized. Future experiments are needed to explicitly test the role of color pattern variation in combining aposematic signals with alternative anti-predatory strategies such as disruptive coloration.

Although we found no direct evidence for an adaptive benefit of variation, within-population pattern heterogeneity could potentially play a significant, yet indirect functional role in polymorphism and polytypism. As aposematic signals are often co-opted *via* sexual selection for mate choice and intra-sexual conflict, patterning may convey information important for intraspecific communication, although research into this possibility has so far predominantly focused on color rather than pattern (Crothers and Cummings, 2013, 2015). The observed variance in pattern could, for example, assist in identifying territory intruders or serve as an honest signal in mate choice or for mediating agonistic interactions (Maan and Cummings, 2008, 2009; Crothers and Cummings, 2013, 2015). In *O. pumilio* future preferences for color-based assortative mating are imprinted during parental care, and to date evidence is limited to coloration because the populations sampled lack pattern elements (Yang et al., 2019). However, were *O. sylvatica* tadpoles to imprint on elements of female phenotypes which include pattern as well as color during maternal trophic egg provisioning, the mother's phenotype could have downstream influences in female mate choice and/or male/male aggression. Albeit indirect, intra-population variance could shape the color pattern evolution in *O. sylvatica* in a mechanism similar to coupled drift proposed in *O. pumilio* by Tazzyman and Iwasa (2010). In coupled drift models female preference (though certainly also potentially applicable to male/male aggression) is linked with color patterns, where preferences diverge according to neutral processes and then reinforce color pattern divergence (Tazzyman and Iwasa, 2010; Gehara et al., 2013). If *O. sylvatica* populations diverge in a method such as coupled drift, then high intra-population variance should enhance between-population divergence, especially those populations which are small or are found in fragmented forest patches.

It has long been predicted that sympatric variance in aposematic signals should be limited, and that which does occur should largely be adaptive (Mappes et al., 2005; Briolat et al., 2019). Recent research adds to a growing body of evidence that this is not universally true, and that relaxed or weak selection can produce or permit significant levels of sympatric intraspecific variation (Richards-Zawacki et al., 2013; Briolat et al., 2019; Bliard et al., 2020). We found that variation within the Perla population of *O. sylvatica* was not divided into visually

distinct categories, and that variation was not associated with particular microhabitat characteristics. Therefore, it is unlikely that signal variance has evolved due to strong selection producing locally adapted phenotypes across a variable fitness landscape, but rather variable patterns remain as purifying selection from predators is weak. These findings highlight how seemingly salient characteristics of animal color may at first evolve through neutral processes, which could have implications for our understanding of the evolution of polymorphism, polytypism, and the evolutionary processes which can ultimately lead to speciation.

DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found below: <https://doi.org/10.5061/dryad.hmgqnk9j0>.

ETHICS STATEMENT

Ethical review and approval was not required for the animal study because animals were only photographed and had no other manipulations. Permits were granted by the Ministerio del Ambiente, Ecuador, for research permission (Permit Number: 014-2019-IC-FLO-DNB/MA).

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AUTHOR CONTRIBUTIONS

JY and JB conceived the manuscript, conducted the analyses, and wrote the manuscript, JY collected the data. Both authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fevo.2022.803996/full#supplementary-material>

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