

Supplementary Information

Ecological and geographical overlap drive plumage evolution and mimicry in woodpeckers

Miller et al.

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Supplementary Table 1 | Principal components analysis of 230,000 CIELAB color values (1,000 samples from each of 230 woodpecker species). The first two PCs explain ~82% of the variance in color. PC1 has high loading values for luminance (L*) and yellowness (positive values of b*), and PC2 has high loading values for greenness (negative values of a*).

	PC1	PC2	PC3
<i>Eigenvalues</i>	1.17	1.04	0.74
<i>Proportion of Variance</i>	0.46	0.36	0.18
<i>Cumulative Proportion of Variance</i>	0.46	0.82	1
Color variable	<i>PC loadings</i>		
L*	0.70	0.29	-0.65
a*	0.01	-0.92	-0.39
b*	0.71	-0.27	0.65

Supplementary Table 2 | Principal components analysis of 138,000 CIELAB back color values (600 samples from each of 230 woodpecker species). The first two PCs explain ~85% of the variance in color. PC1 has high loading values for luminance (L*) and yellowness (positive values of b*), and PC2 has high loading values for greenness (negative values of a*).

	PC1	PC2	PC3
<i>Eigenvalues</i>	1.27	0.96	0.68
<i>Proportion of Variance</i>	0.54	0.31	0.15
<i>Cumulative Proportion of Variance</i>	0.54	0.85	1.00
Color variable	<i>PC loadings</i>		
L*	0.67	-0.21	-0.71
a*	-0.32	-0.95	-0.03
b*	0.67	-0.25	0.70

Supplementary Table 3 | Principal components analysis of 46,000 CIELAB head color values (200 samples from each of 230 woodpecker species). The first two PCs explain ~87% of the variance in color. PC1 has high loading values for greenness (negative values of a*) and blueness (negative values of b*), and PC2 has high loading values for luminance (L*).

	PC1	PC2	PC3
<i>Eigenvalues</i>	1.19	1.09	0.63
<i>Proportion of Variance</i>	0.47	0.39	0.13
<i>Cumulative Proportion of Variance</i>	0.47	0.87	1.00
Color variable	<i>PC loadings</i>		
L*	0.02	0.88	0.48
a*	-0.71	-0.33	0.62
b*	-0.70	0.35	-0.62

Supplementary Table 4 | Principal components analysis of 23,000 CIELAB belly color values (100 samples from each of 230 woodpecker species). The first two PCs explain ~81% of the variance in color. PC1 has high loading values for luminance (L*) and yellowness (positive values of b*), and PC2 has high loading values for greenness (negative values of a*).

	PC1	PC2	PC3
<i>Eigenvalues</i>	1.12	1.08	0.75
<i>Proportion of Variance</i>	0.42	0.39	0.19
<i>Cumulative Proportion of Variance</i>	0.42	0.81	1.00
Color variable	<i>PC loadings</i>		
L*	0.70	-0.39	0.60
a*	0.02	0.85	0.53
b*	0.72	0.35	-0.60

Supplementary Table 5 | Principal components analysis of pattern energy spectra across 230 species of woodpeckers. The three PC axes explain ~93% of the variance in pattern across all species.

	PC1	PC2	PC3
Standard deviation	3.08	2.12	1.34
Proportion of Variance	0.56	0.26	0.11
Cumulative Proportion	0.56	0.82	0.93

Supplementary Table 6 | Principal components analysis of pattern energy spectra across the backs of 230 species of woodpeckers. The three PC axes explain ~93% of the variance in pattern across all species.

	PC1	PC2	PC3
Standard deviation	2.85	2.46	1.29
Proportion of Variance	0.48	0.36	0.10
Cumulative Proportion	0.48	0.84	0.93

Supplementary Table 7 | Principal components analysis of pattern energy spectra across the heads of 230 species of woodpeckers. The three PC axes explain ~86% of the variance in pattern across all species.

	PC1	PC2	PC3
Standard deviation	2.80	1.91	1.75
Proportion of Variance	0.46	0.21	0.18
Cumulative Proportion	0.46	0.67	0.86

Supplementary Table 8 | Principal components analysis of pattern energy spectra across the bellies of 230 species of woodpeckers. The three PC axes explain ~92% of the variance in pattern across all species.

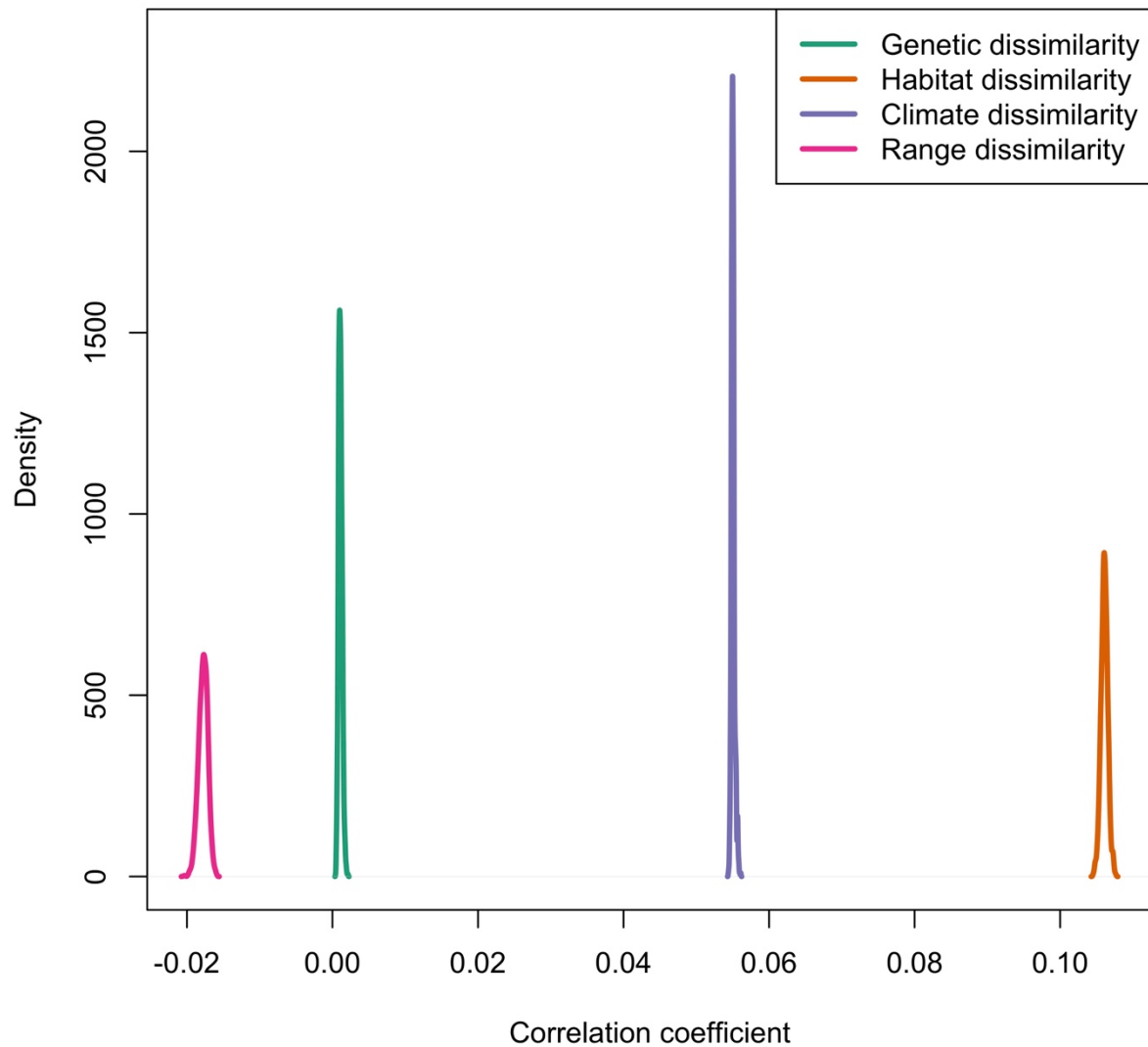
	PC1	PC2	PC3
Standard deviation	2.84	2.45	1.26
Proportion of Variance	0.48	0.35	0.09
Cumulative Proportion	0.48	0.83	0.92

Supplementary Table 9 | Specimens from the Cornell University Museum of Vertebrates (CUMV) used for color and pattern analyses.

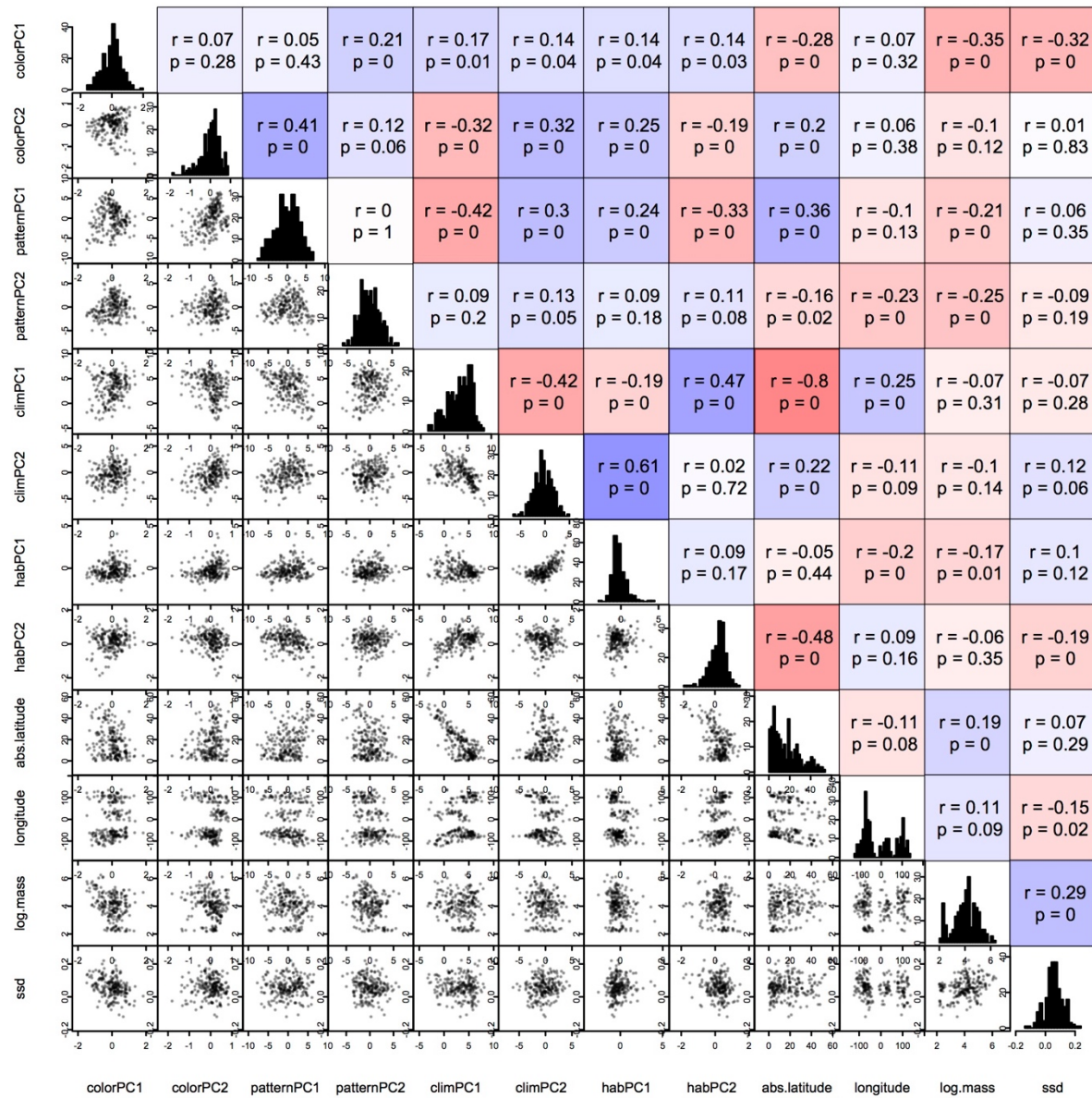
Species (our taxonomy)	Species (on CUMV tag)	CUMV tag subspecies	CUMV collection number	Location
<i>Xiphidiopicus percussus</i>	<i>Xiphidiopicus percussus</i>	<i>percussus</i>	22746	Cuba
<i>Xiphidiopicus percussus</i>	<i>Xiphidiopicus percussus</i>	<i>percussus</i>	6498	Cuba
<i>Xiphidiopicus percussus</i>	<i>Xiphidiopicus percussus</i>	<i>percussus</i>	6497	Cuba
<i>Sphyrapicus varius</i>	<i>Sphyrapicus varius</i>	NA	55875	NY
<i>Sphyrapicus varius</i>	<i>Sphyrapicus varius</i>	NA	53166	NY
<i>Sphyrapicus varius</i>	<i>Sphyrapicus varius</i>	NA	30185	NY
<i>Melanerpes carolinus</i>	<i>Melanerpes carolinus</i>	NA	51400	NY
<i>Melanerpes carolinus</i>	<i>Melanerpes carolinus</i>	NA	55391	NY
<i>Melanerpes carolinus</i>	<i>Melanerpes carolinus</i>	NA	6198	TN
<i>Dendrocopos major</i>	<i>Dendrocopos major</i>	NA	6542	France
<i>Dendrocopos major</i>	<i>Dendrocopos major</i>	NA	6540	Germany
<i>Dendrocopos major</i>	<i>Dendrocopos major</i>	NA	6541	Italy
<i>Colaptes rubiginosus</i>	<i>Colaptes rubiginosus</i>	NA	34519	Chiapas
<i>Colaptes rubiginosus</i>	<i>Colaptes rubiginosus</i>	NA	6335	Motzorongo, Veracruz
<i>Mesopicos goertae</i>	<i>Mesopicos goertae</i>	NA	33636	Uganda
<i>Mesopicos goertae</i>	<i>Dendropicos spodocephalus</i>	NA	33640	Kenya
<i>Mesopicos goertae</i>	<i>Dendropicos spodocephalus</i>	<i>rhodeogaster</i>	6524	illegible
<i>Picoides villosus</i>	<i>Picoides villosus</i>	<i>villosus</i>	6406	NY
<i>Picoides villosus</i>	<i>Picoides villosus</i>	<i>villosus</i>	6407	NY
<i>Picoides villosus</i>	<i>Picoides villosus</i>	<i>villosus</i>	6415	NY
<i>Picoides albolarvatus</i>	<i>Picoides albolarvatus</i>	<i>albolarvatus</i>	6736	CA
<i>Picoides albolarvatus</i>	<i>Picoides albolarvatus</i>	<i>albolarvatus</i>	6733	CA
<i>Picoides albolarvatus</i>	<i>Picoides albolarvatus</i>	NA	22183	WA
<i>Picus viridis</i>	<i>Picus viridis</i>	<i>virescens</i>	7762	England
<i>Picus viridis</i>	<i>Picus viridis</i>	<i>virescens</i>	7763	England
<i>Dendropicos fuscescens</i>	<i>Dendropicos fuscescens</i>	<i>fuscescens</i>	6530	Angola
<i>Dendropicos fuscescens</i>	<i>Dendropicos fuscescens</i>	<i>fuscescens</i>	6529	Angola
<i>Dendropicos fuscescens</i>	<i>Dendropicos fuscescens</i>	NA	32433	Southwest Africa
<i>Picoides pubescens</i>	<i>Picoides pubescens</i>	<i>medianus</i>	6569	NY
<i>Picoides pubescens</i>	<i>Picoides pubescens</i>	<i>medianus</i>	6593	NY
<i>Picoides pubescens</i>	<i>Picoides pubescens</i>	<i>medianus</i>	6571	NY
<i>Veniliornis kirkii</i>	<i>Veniliornis kirkii</i>	NA	26606	Colombia
<i>Veniliornis kirkii</i>	<i>Veniliornis kirkii</i>	<i>neglectus</i>	6518	Costa Rica

<i>Dryocopus pileatus</i>	<i>Dryocopus pileatus</i>	NA	52879	NY
<i>Dryocopus pileatus</i>	<i>Dryocopus pileatus</i>	NA	54931	NY
<i>Dryocopus pileatus</i>	<i>Dryocopus pileatus</i>	NA	53082	NY
<i>Dinopium benghalense</i>	<i>Dinopium benghalense</i>	<i>benghalense</i>	23454	India
<i>Campephilus</i>	<i>Campephilus</i>	NA	29193	Honduras
<i>guatemalensis</i>	<i>guatemalensis</i>			
<i>Campephilus</i>	<i>Campephilus</i>	<i>guatemalensis</i>	6397	Costa Rica
<i>guatemalensis</i>	<i>guatemalensis</i>			
<i>Campephilus</i>	<i>Campephilus</i>	<i>regius</i>	6402	Mexico
<i>guatemalensis</i>	<i>guatemalensis</i>			
<i>Campethera nivos</i>	<i>Campethera nivos</i>	<i>efulenensis</i>	6519	Cameroon
<i>Campethera nivos</i>	<i>Campethera nivos</i>	NA	29775	Nigeria
<i>Campethera nivos</i>	<i>Campethera nivos</i>	NA	33632	Uganda
<i>Gecinulus grantia</i>	<i>Gecinulus grantia</i>	NA	6464	India
<i>Blythipicus pyrrhotis</i>	<i>Blythipicus pyrrhotis</i>	NA	6479	India
<i>Hemicircus canente</i>	<i>Hemicircus canente</i>	<i>canente</i>	6501	India
<i>Chrysocolaptes</i>	<i>Chrysocolaptes</i>	NA	6486	illegible
<i>guttacristatus</i>	<i>guttacristatus</i>			
<i>Chrysocolaptes</i>	<i>Chrysocolaptes</i>	NA	23486	Burma
<i>guttacristatus</i>	<i>guttacristatus</i>			
<i>Colaptes auratus</i>	<i>Colaptes auratus</i>	NA	48699	NY
<i>Colaptes auratus</i>	<i>Colaptes auratus</i>	NA	28168	NY
<i>Colaptes auratus</i>	<i>Colaptes auratus</i>	NA	14975	NY
<i>Colaptes auratus</i>	<i>Colaptes auratus</i>	NA	48717	NY
<i>Meiglyptes tristis</i>	<i>Meiglyptes tristis</i>	<i>grammithorax</i>	6466	India
<i>Celeus castaneus</i>	<i>Celeus castaneus</i>	NA	6471	Mexico
<i>Celeus castaneus</i>	<i>Celeus castaneus</i>	NA	6473	Costa Rica
<i>Celeus castaneus</i>	<i>Celeus castaneus</i>	NA	6472	Costa Rica

Accounting for phylogenetic uncertainty in the multiple distance matrix regression

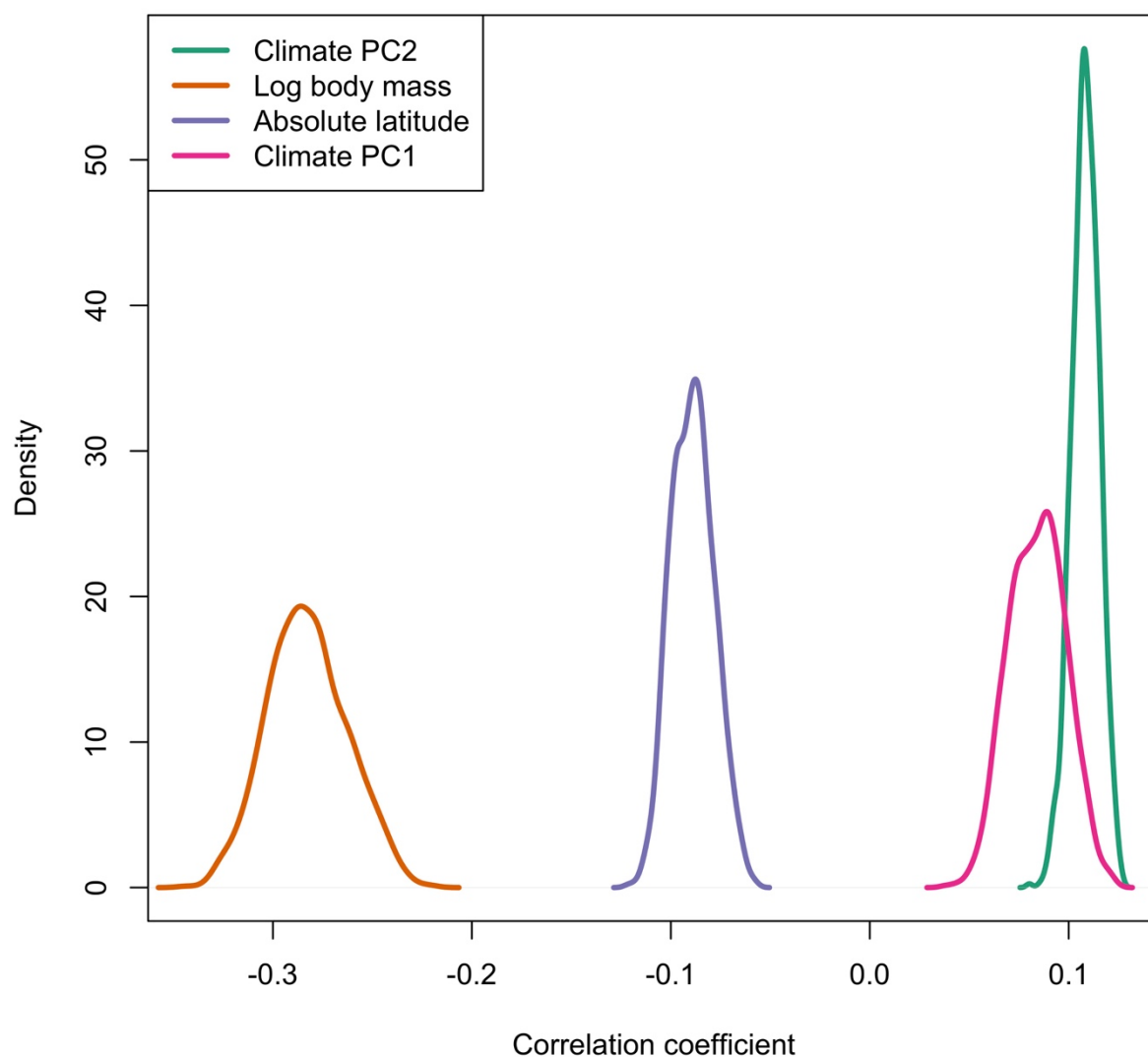


Supplementary Fig. 1 | Correlations between plumage and each of the independent variables are not strongly influenced by phylogenetic uncertainty. Correlation coefficients between the independent matrices (genetic, habitat, and climate dissimilarity) and the dependent matrix (plumage dissimilarity) have narrow distributions (kernel density estimates shown).



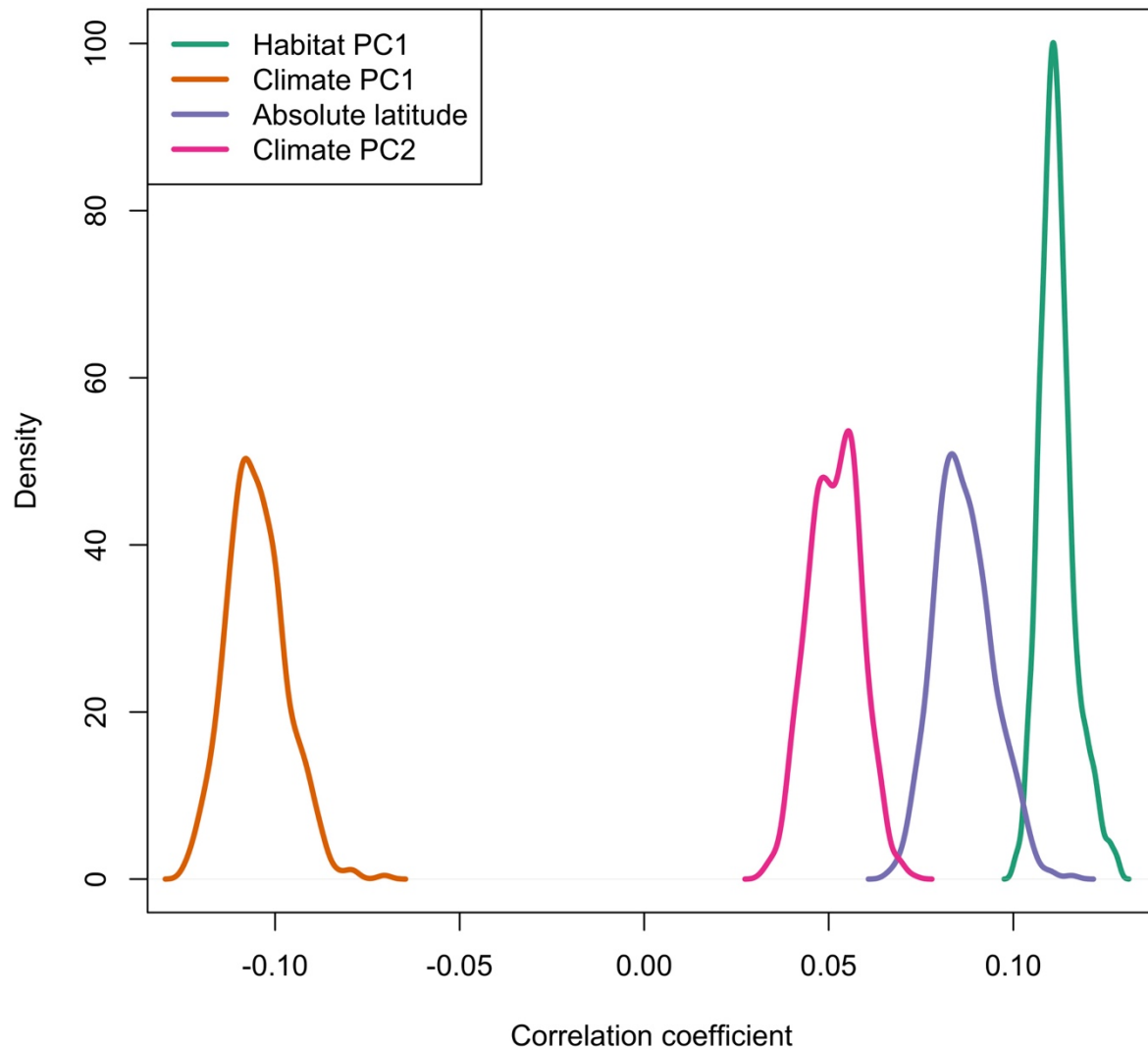
Supplementary Fig. 2 | Paneled correlation plot showing the relationships between all species-level averaged traits. Climate, habitat, latitude, and woodpecker body mass and plumage appearance exhibit a variety of intercorrelations. The lower triangle panels show the pairwise trait relationships. The diagonal panels contain histograms of the indicated variable. The upper triangle panels summarize the Pearson's correlation coefficient and the significance of the relationship, both numerically and in color, where bright red indicates a strong negative correlation, and bright blue indicates a strong positive correlation.

Accounting for phylogenetic uncertainty in colorPC1 PGLS



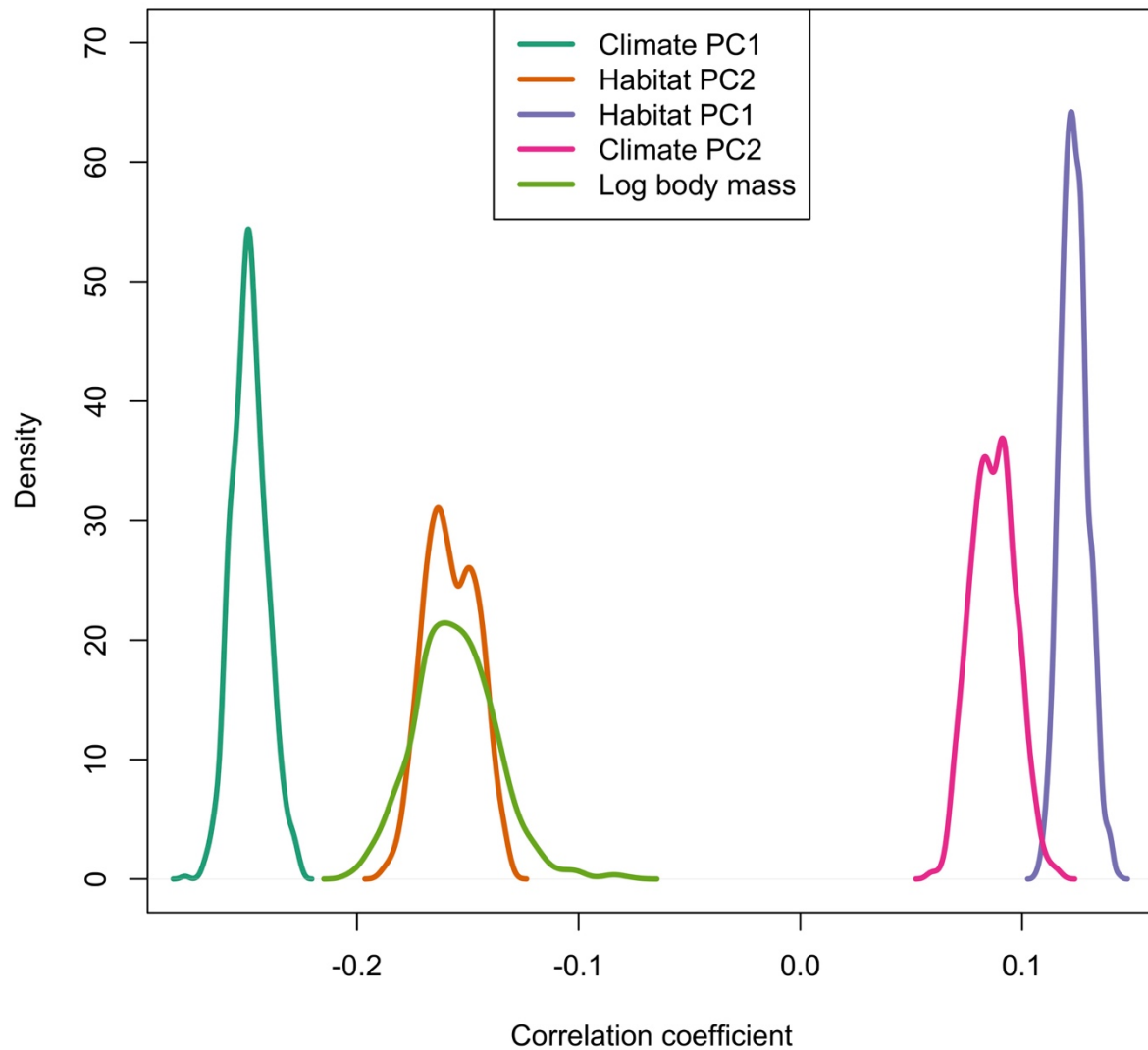
Supplementary Fig. 3 | Correlations between colorPC1 and each of the independent variables are not strongly influenced by phylogenetic uncertainty. Correlation coefficients between the independent variables and colorPC1 have narrow distributions (kernel density estimates shown).

Accounting for phylogenetic uncertainty in colorPC2 PGLS



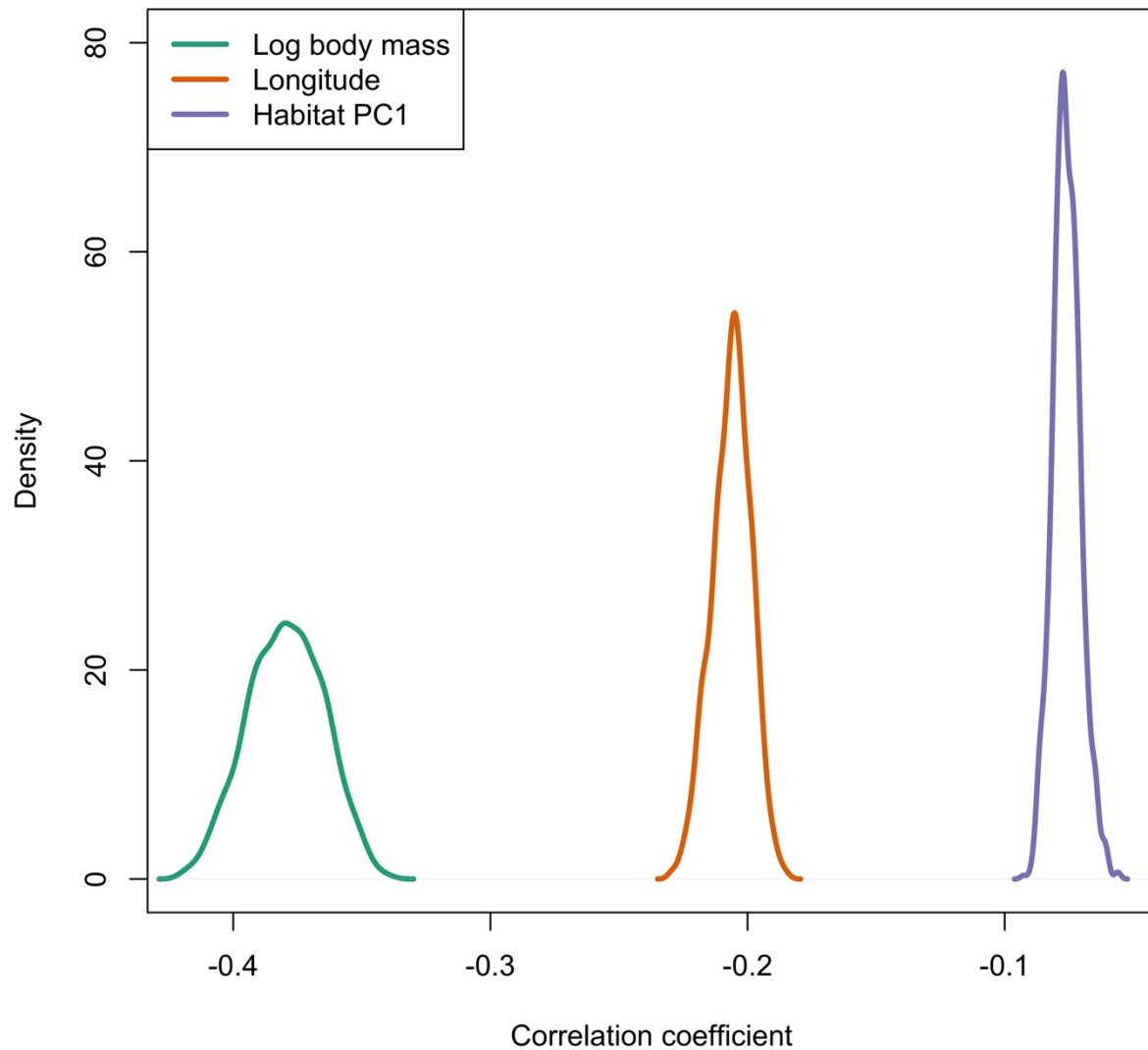
Supplementary Fig. 4 | Correlations between colorPC1 and each of the independent variables are not strongly influenced by phylogenetic uncertainty. Correlation coefficients between the independent variables and colorPC2 have narrow distributions (kernel density estimates shown).

Accounting for phylogenetic uncertainty in patternPC1 PGLS

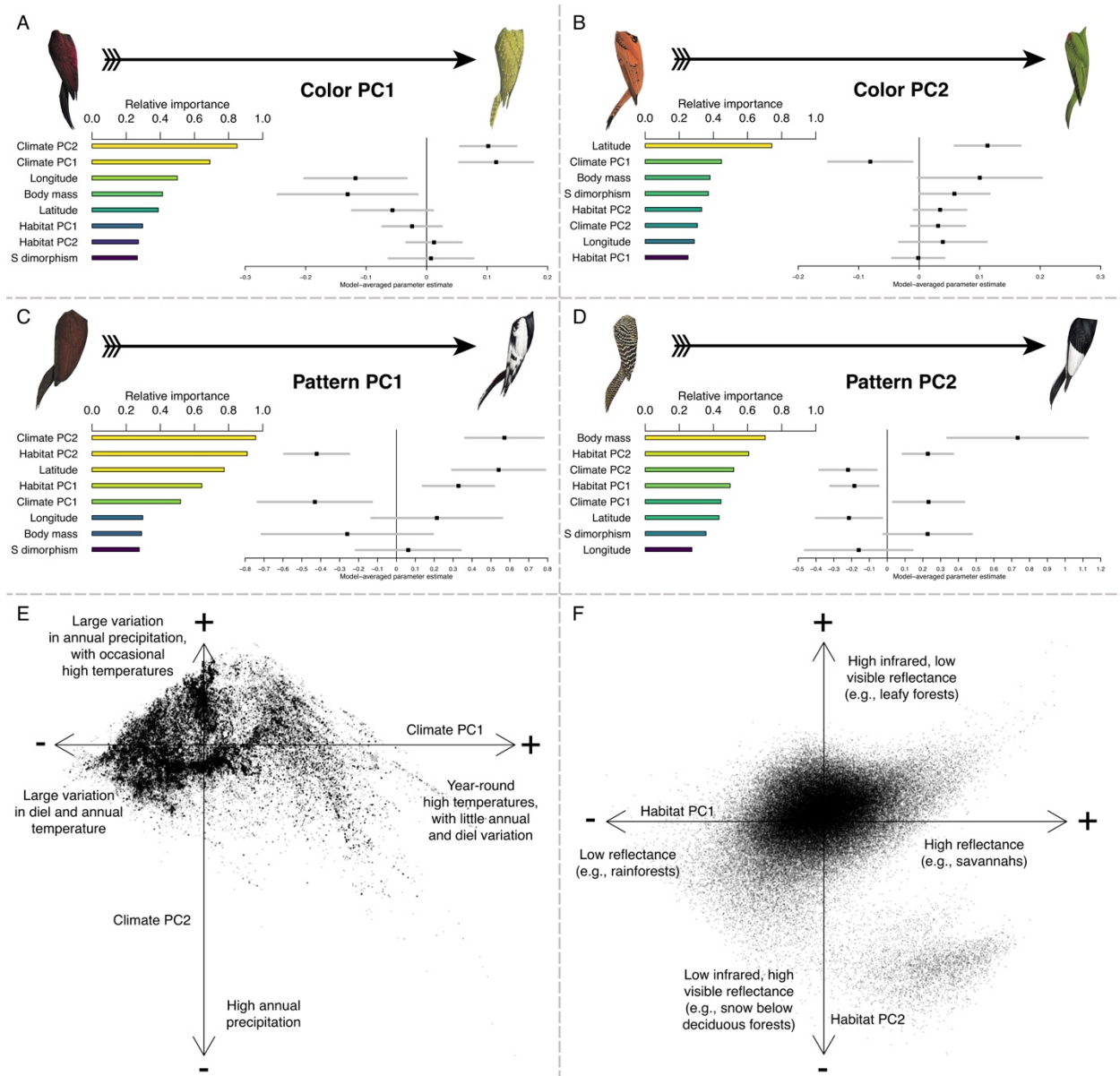


Supplementary Fig. 5 | Correlations between colorPC1 and each of the independent variables are not strongly influenced by phylogenetic uncertainty. Correlation coefficients between the independent variables and patternPC1 have narrow distributions (kernel density estimates shown).

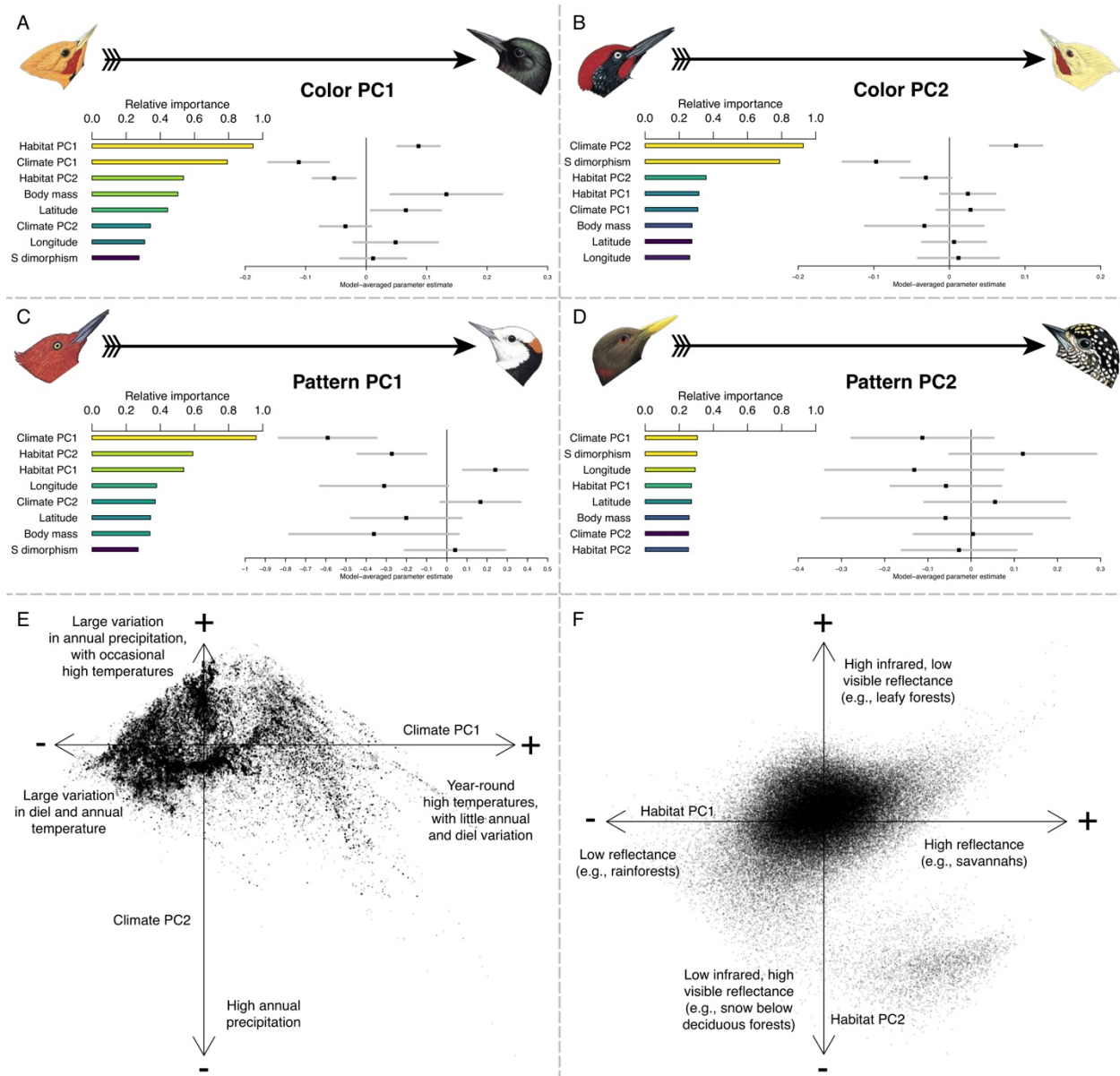
Accounting for phylogenetic uncertainty in patternPC2 PGLS



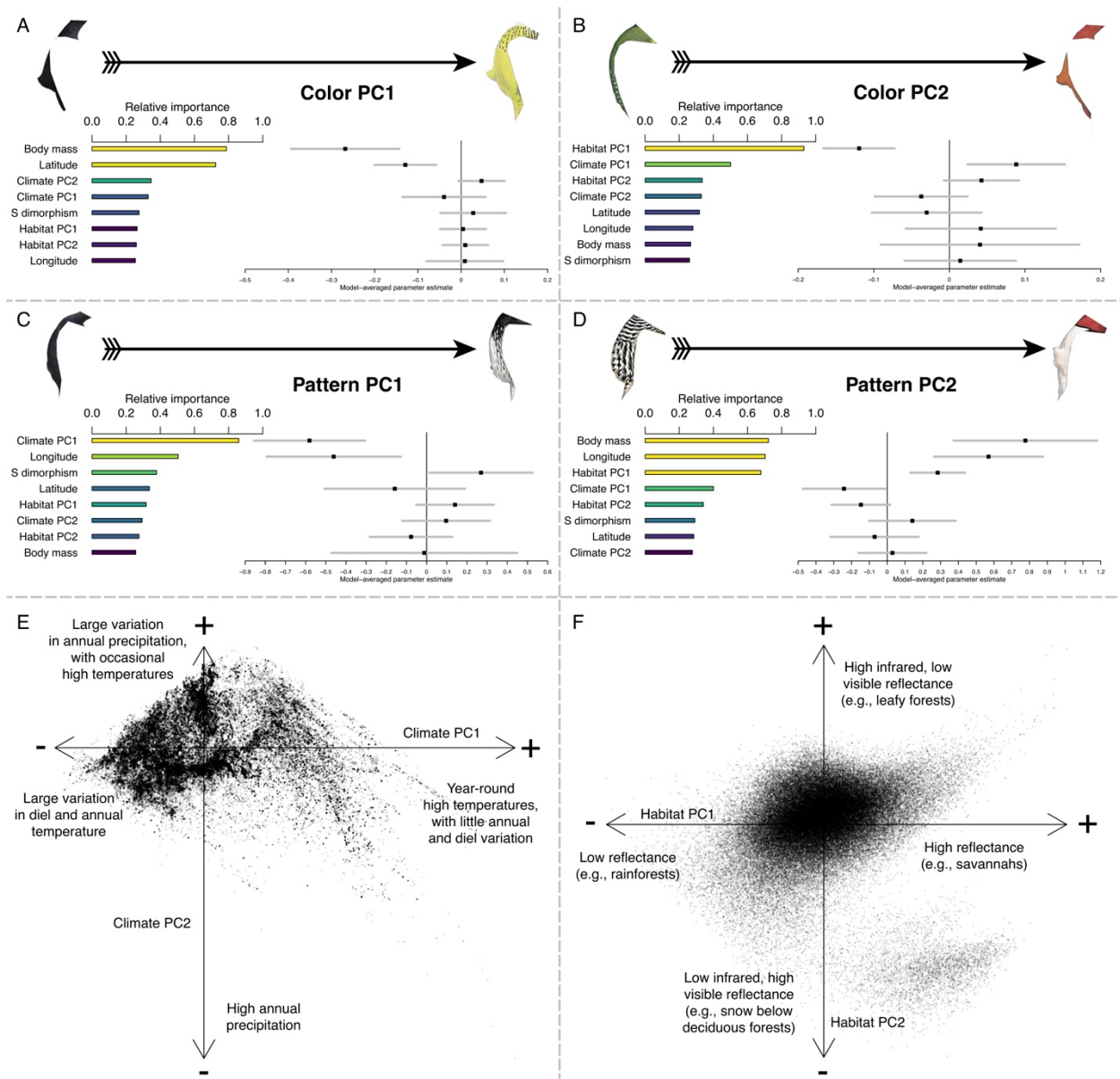
Supplementary Fig. 6 | Correlations between colorPC1 and each of the independent variables are not strongly influenced by phylogenetic uncertainty. Correlation coefficients between the independent variables and patternPC2 have narrow distributions (kernel density estimates shown).



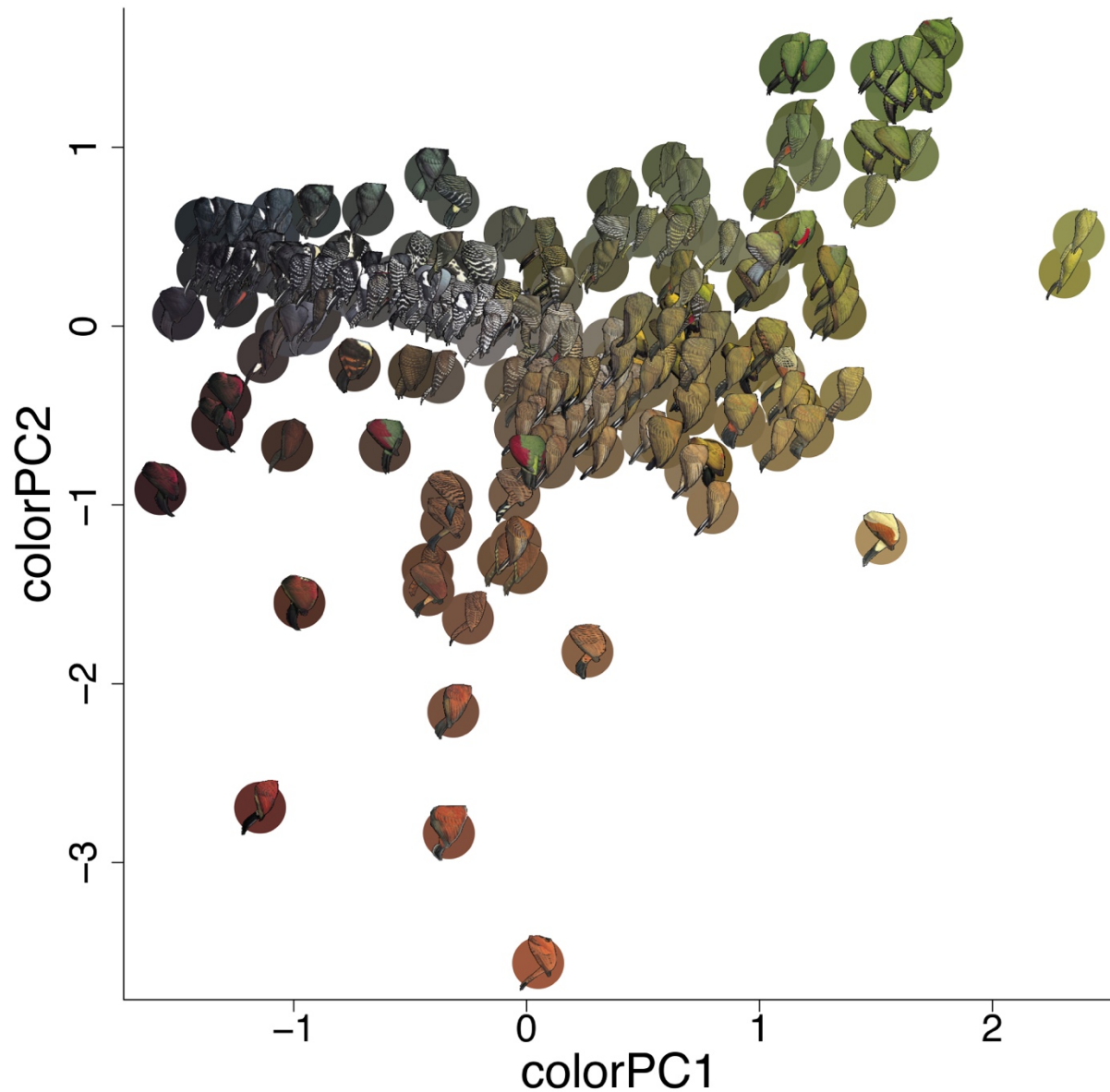
Supplementary Fig. 7 | Back-specific variable importance scores and model-averaged parameter estimates from phylogenetic generalized least squares regressions. These quantify how colour and pattern vary as a function of climate, habitat, body mass, sexual size dimorphism, latitude and longitude, with summaries of the climate and habitat principal component analyses (PCA). Model-averaged p -values of explanatory factors are colour-coded from yellow to blue; only factors with p -values < 0.05 are coloured yellow and discussed here. (a) Dark birds are heavier and occur in wetter climates. (b) Greenish (as opposed to reddish) birds are found in more open habitats. (c) Less-patterned birds are found in aseasonal climates, open habitats, and temperate forests. (d) Birds patterned in large plumage elements, such as large colour patches, tend to be larger in body size. (e) Climate PCA results, illustrating the distribution of woodpeckers in climate space, with qualitative descriptions of the first two PC axes. (f) Habitat PCA results, showing the distribution of woodpeckers across global habitats, with qualitative descriptions of the first two PC axes. Illustrations © HBW Alive/Lynx Edicions.



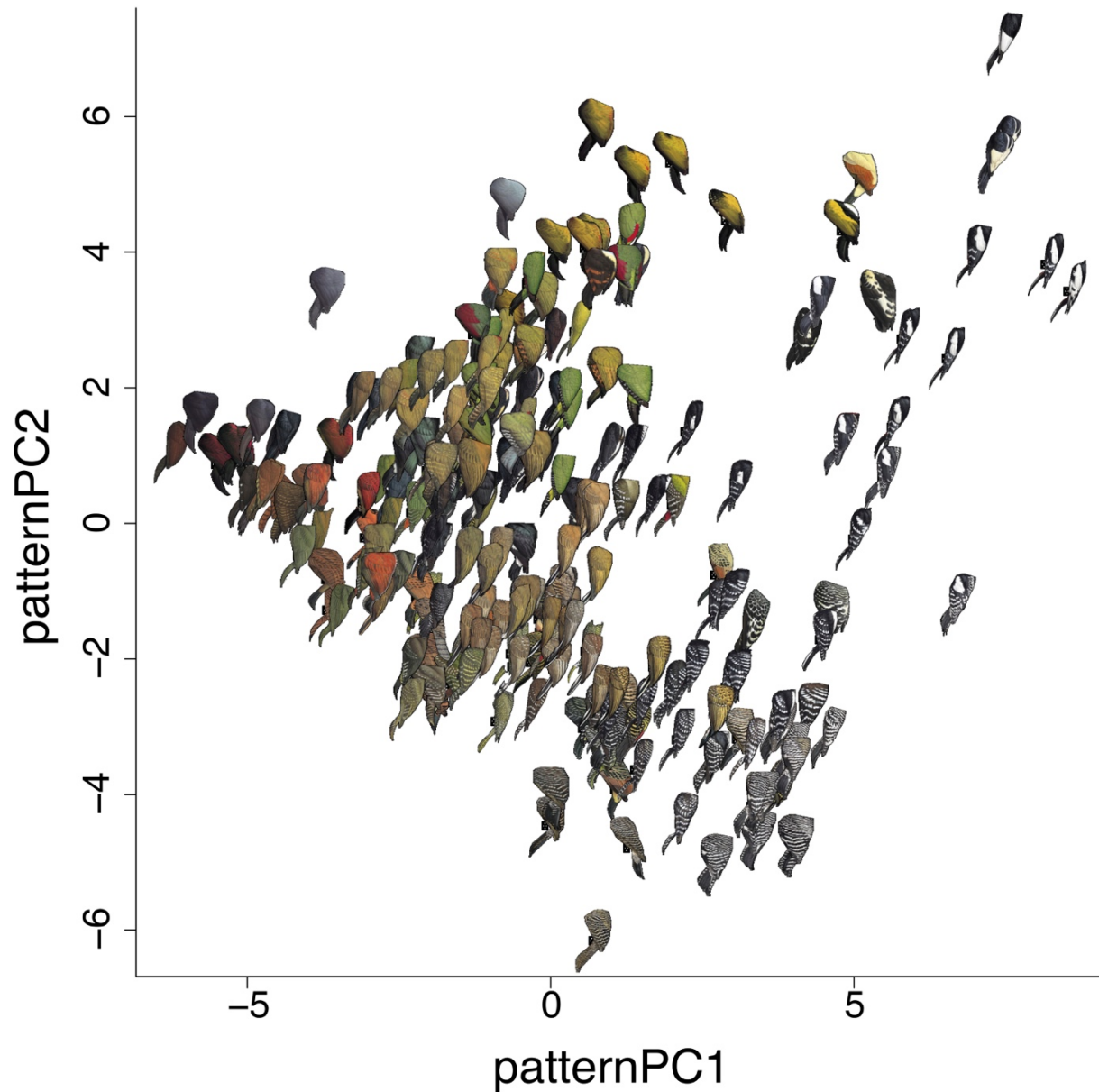
Supplementary Fig. 8 | Head-specific variable importance scores and model-averaged parameter estimates from phylogenetic generalized least squares regressions. These quantify how colour and pattern vary as a function of climate, habitat, body mass, sexual size dimorphism, latitude and longitude, with summaries of the climate and habitat principal component analyses (PCA). Model-averaged p -values of explanatory factors are colour-coded from yellow to blue; only factors with p -values < 0.05 are coloured yellow and discussed here. (a) Dark birds are heavier and occur in wetter climates. (b) Greenish (as opposed to reddish) birds are found in more open habitats. (c) Less-patterned birds are found in aseasonal climates, open habitats, and temperate forests. (d) Birds patterned in large plumage elements, such as large colour patches, tend to be larger in body size. (e) Climate PCA results, illustrating the distribution of woodpeckers in climate space, with qualitative descriptions of the first two PC axes. (f) Habitat PCA results, showing the distribution of woodpeckers across global habitats, with qualitative descriptions of the first two PC axes. Illustrations © HBW Alive/Lynx Edicions.



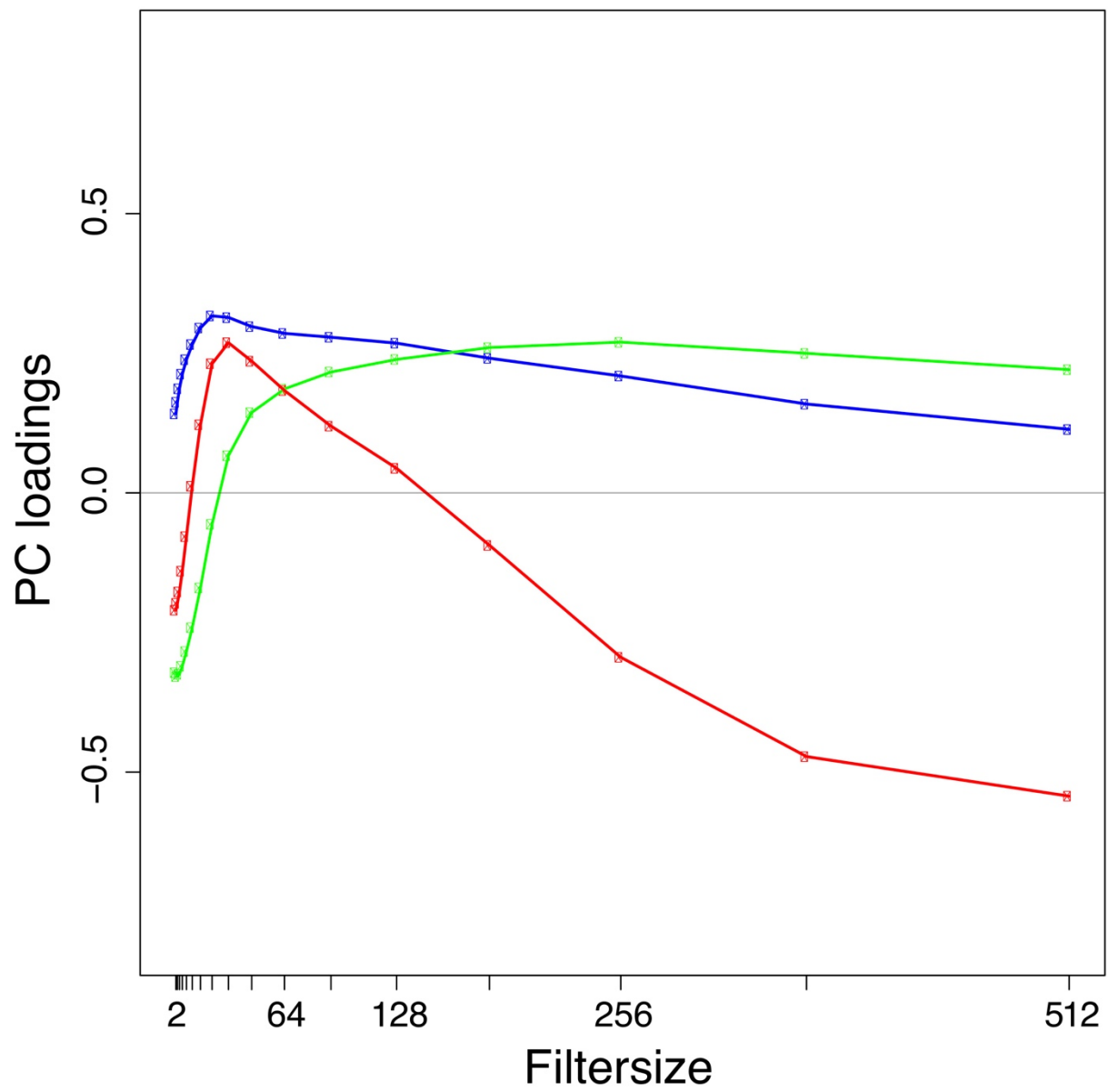
Supplementary Fig. 9 | Belly-specific variable importance scores and model-averaged parameter estimates from phylogenetic generalized least squares regressions. These quantify how colour and pattern vary as a function of climate, habitat, body mass, sexual size dimorphism, latitude and longitude, with summaries of the climate and habitat principal component analyses (PCA). Model-averaged p -values of explanatory factors are colour-coded from yellow to blue; only factors with p -values < 0.05 are coloured yellow and discussed here. (a) Dark birds are heavier and occur in wetter climates. (b) Greenish (as opposed to reddish) birds are found in more open habitats. (c) Less-patterned birds are found in aseasonal climates, open habitats, and temperate forests. (d) Birds patterned in large plumage elements, such as large colour patches, tend to be larger in body size. (e) Climate PCA results, illustrating the distribution of woodpeckers in climate space, with qualitative descriptions of the first two PC axes. (f) Habitat PCA results, showing the distribution of woodpeckers across global habitats, with qualitative descriptions of the first two PC axes. Illustrations © HBW Alive/Lynx Edicions.



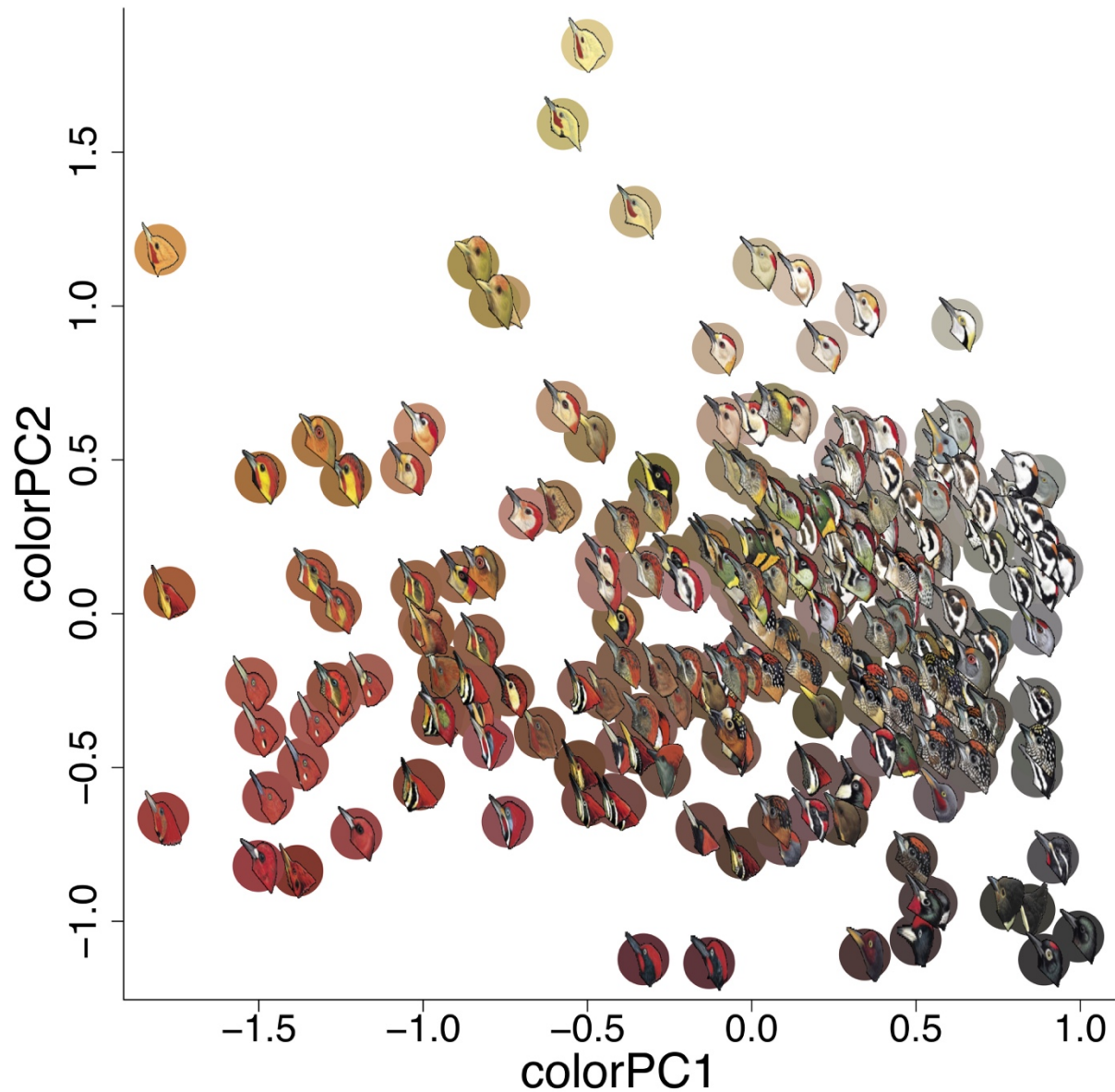
Supplementary Fig. 10 | Back-specific principal components analysis (PCA) of species-averaged woodpecker colour values. Principal component one (colourPC1) explains 45% of the variation in measured colour scores. Higher PC1 scores correspond to greater luminance values, and more yellow and less blue. Principal component two (colourPC2) explains an additional 36% of variation in overall colour scores. Higher PC2 scores correspond to more green and less red colouration. Coloured circles behind each woodpecker species correspond to the average CIE L*a*b scores for the 600 randomly selected colour samples from that species' back. Illustrations © HBW Alive/Lynx Edicions.



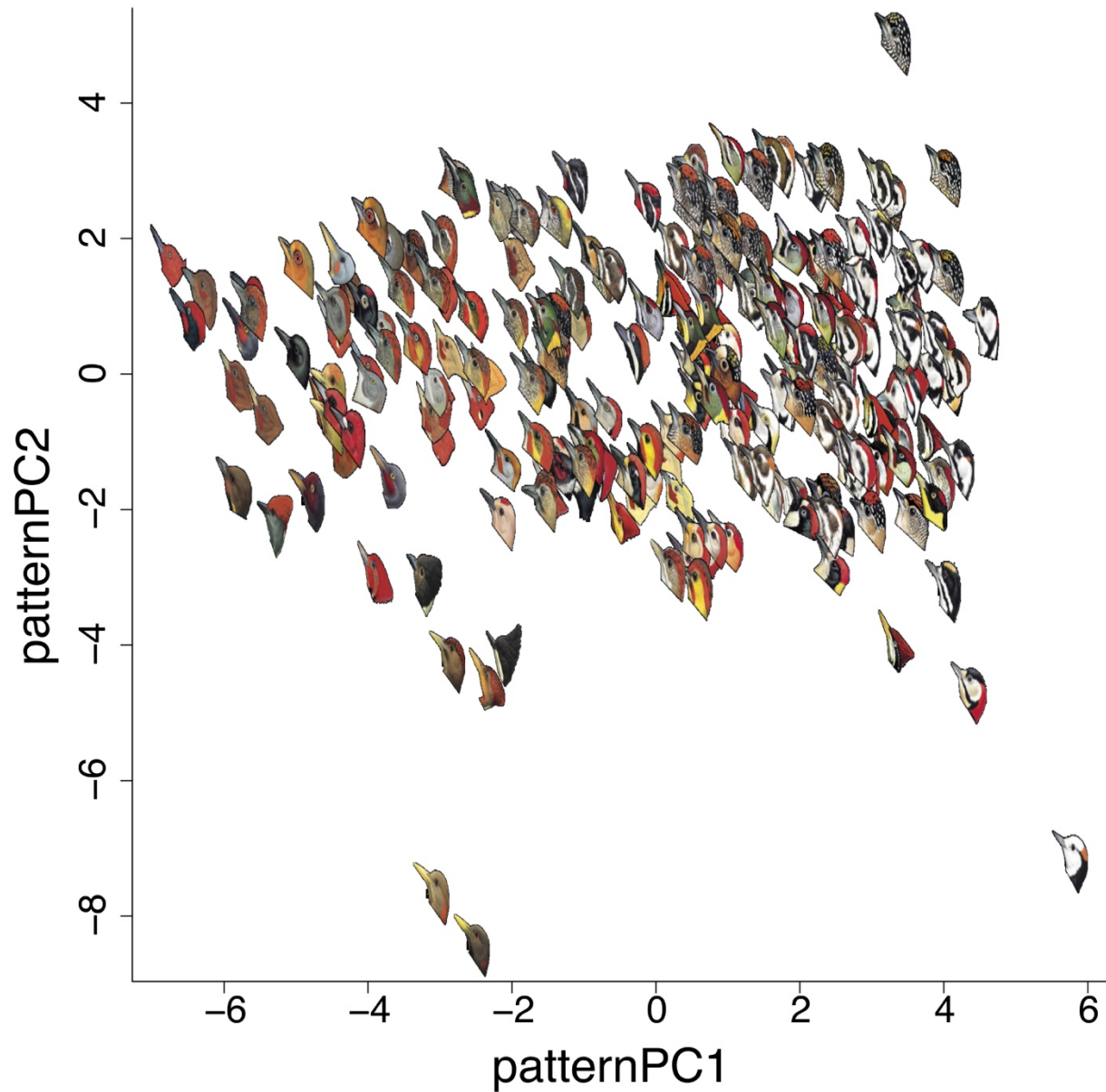
Supplementary Fig. 11 | Back-specific major axes of plumage pattern variation quantified using granularity analysis, then summarized with a principal components analysis (PCA) for species-level approaches. Pattern PC1 and PC2, collectively, account for 82.1% of variation across woodpeckers. Exemplar species (i-iv) illustrate extreme variation in PC1: (i) exhibits high energy scores across most pattern element sizes, with small, medium, and large pattern elements; (ii) has low energy scores across the spectrum, with few pattern elements of any size. And, extreme variation in PC2: (iii) has many small pattern elements and few of any other sizes; (iv) has only medium and large size pattern elements. Illustrations © HBW Alive/Lynx Edicions.



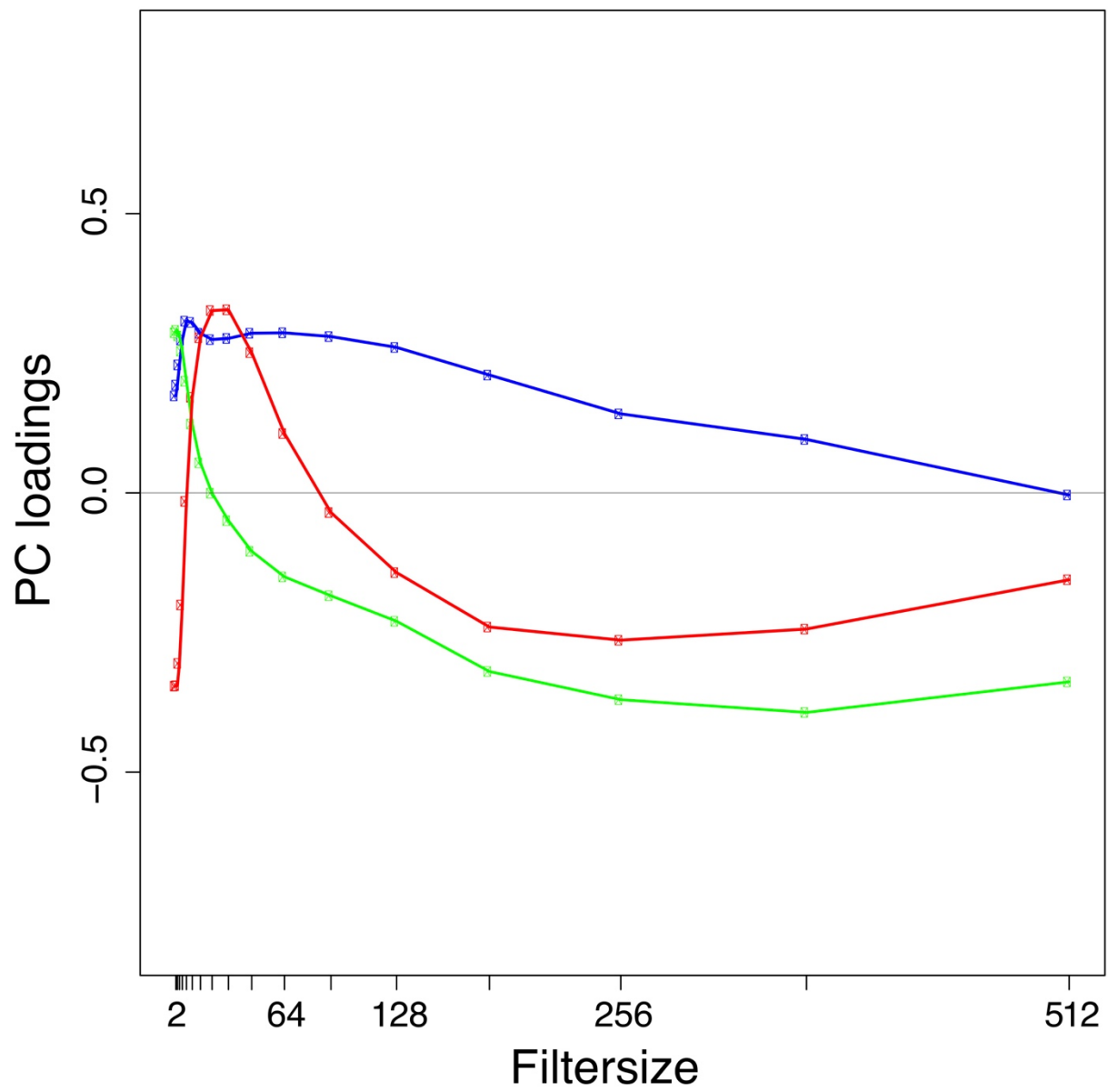
Supplementary Fig. 12 | Principal component loadings of back pattern. Principal component loadings of pattern energy spectra across filter size for all species reveals how back pattern elements of different sizes influence PC scores.



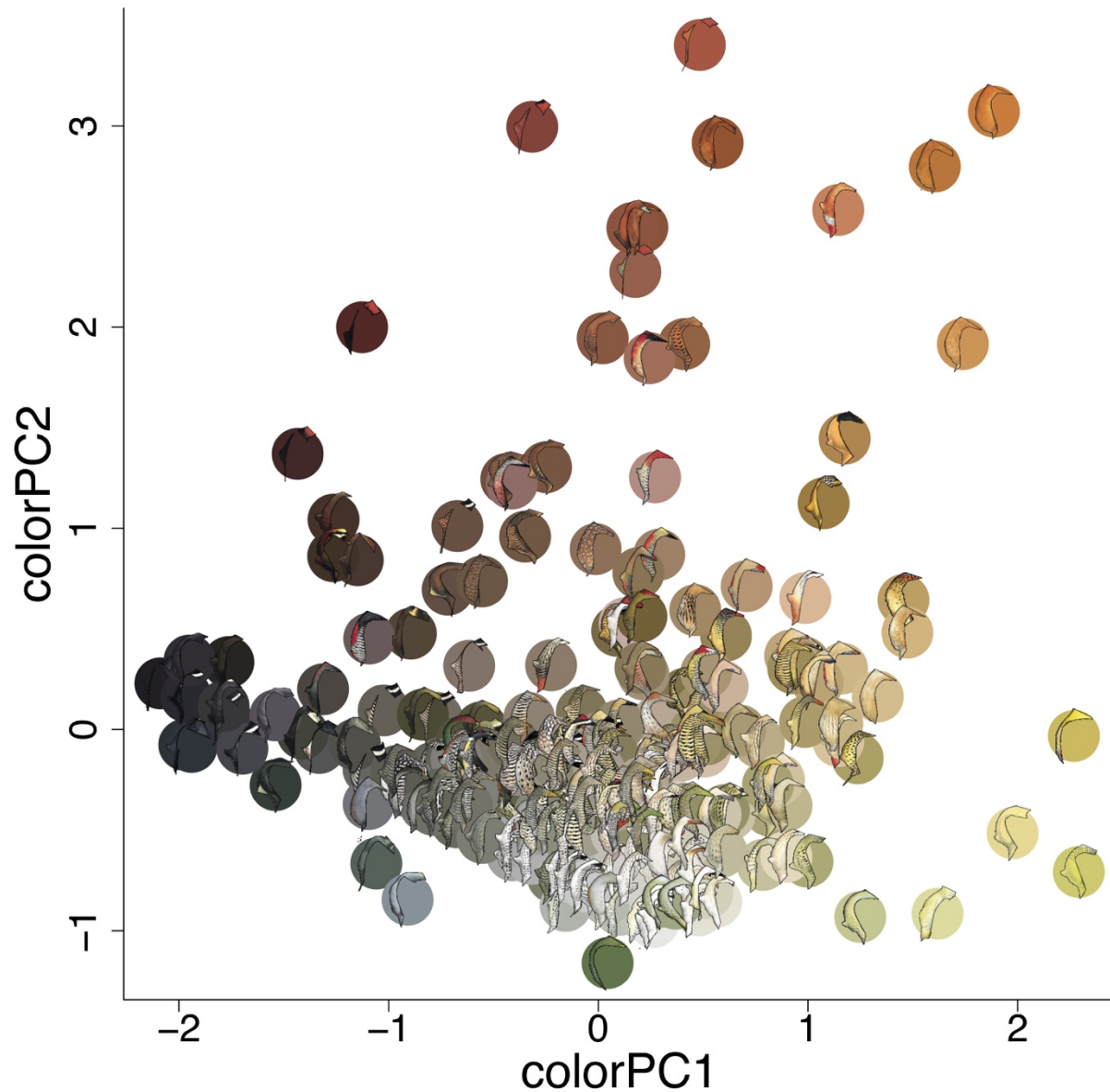
Supplementary Fig. 13 | Head-specific principal components analysis (PCA) of species-averaged woodpecker colour values. Principal component one (colourPC1) explains 45% of the variation in measured colour scores. Higher PC1 scores correspond to greater luminance values, and more yellow and less blue. Principal component two (colourPC2) explains an additional 36% of variation in overall colour scores. Higher PC2 scores correspond to more green and less red colouration. Coloured circles behind each woodpecker species correspond to the average CIE L*a*b scores for the 200 randomly selected colour samples from that species' head. Illustrations © HBW Alive/Lynx Edicions.



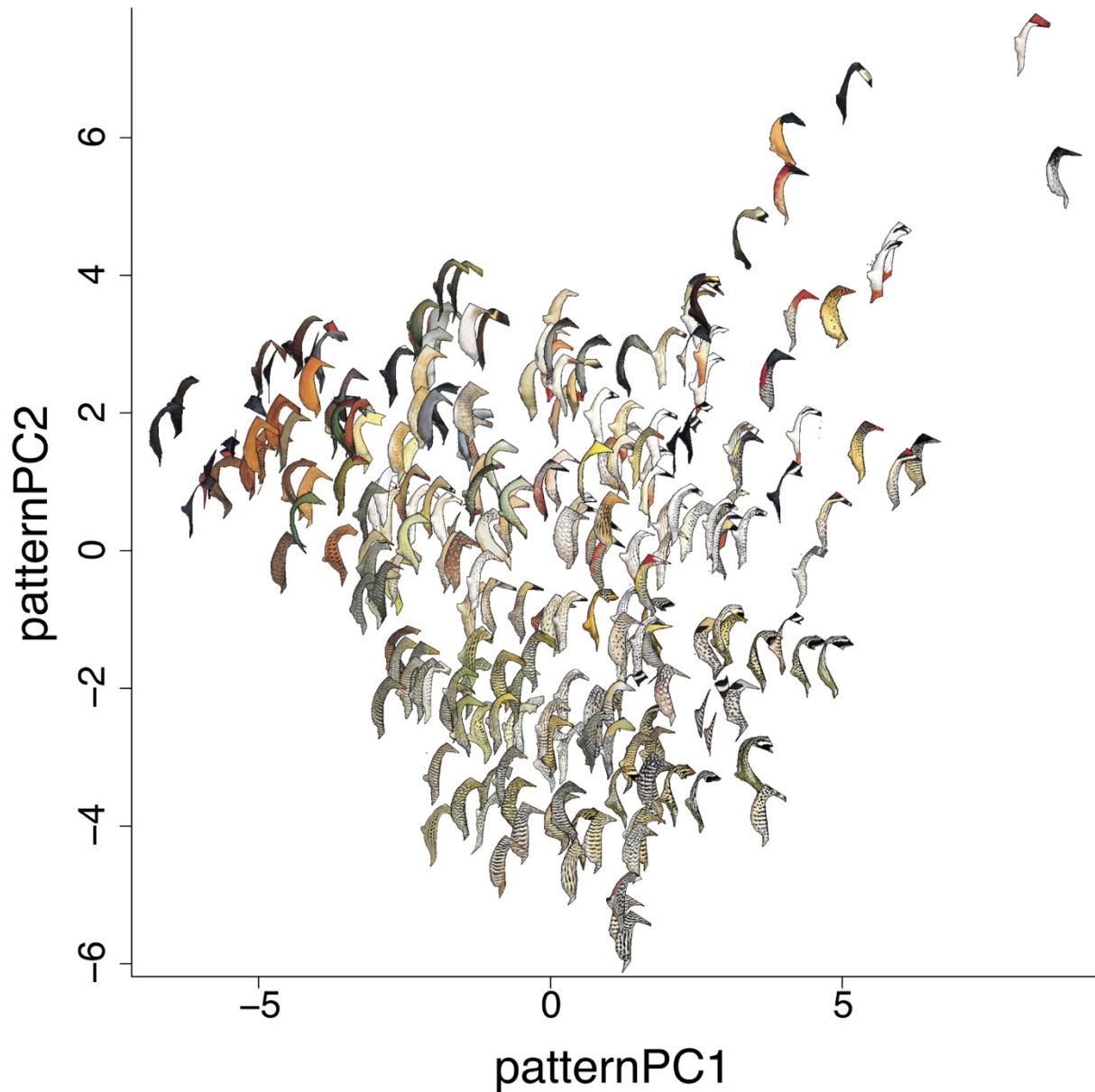
Supplementary Fig. 14 | Head-specific major axes of plumage pattern variation quantified using granularity analysis, then summarized with a principal components analysis (PCA) for species-level approaches. Pattern PC1 and PC2, collectively, account for 82.1% of variation across woodpeckers. Exemplar species (i-iv) illustrate extreme variation in PC1: (i) exhibits high energy scores across most pattern element sizes, with small, medium, and large pattern elements; (ii) has low energy scores across the spectrum, with few pattern elements of any size. And, extreme variation in PC2: (iii) has many small pattern elements and few of any other sizes; (iv) has only medium and large size pattern elements. Illustrations © HBW Alive/Lynx Edicions.



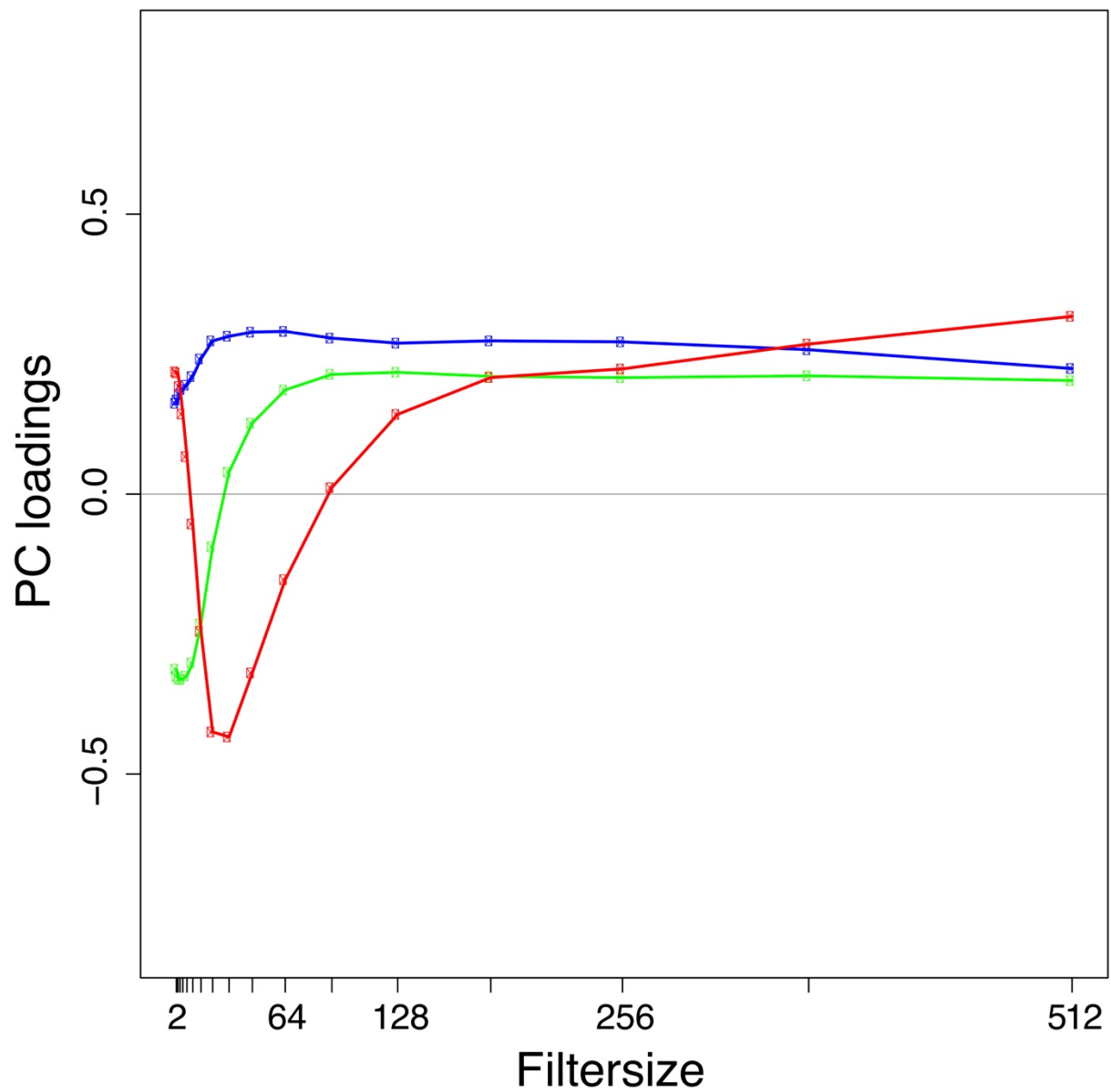
Supplementary Fig. 15 | Principal component loadings of head pattern. Principal component loadings of pattern energy spectra across filter size for all species reveals how head pattern elements of different sizes influence PC scores.



Supplementary Fig. 16 | Belly-specific principal components analysis (PCA) of species-averaged woodpecker colour values. Principal component one (colourPC1) explains 45% of the variation in measured colour scores. Higher PC1 scores correspond to greater luminance values, and more yellow and less blue. Principal component two (colourPC2) explains an additional 36% of variation in overall colour scores. Higher PC2 scores correspond to more green and less red colouration. Coloured circles behind each woodpecker species correspond to the average CIE L*a*b scores for the 100 randomly selected colour samples from that species' belly. Illustrations © HBW Alive/Lynx Edicions.



Supplementary Fig. 17 | Belly-specific major axes of plumage pattern variation quantified using granularity analysis, then summarized with a principal components analysis (PCA) for species-level approaches. Pattern PC1 and PC2, collectively, account for 82.1% of variation across woodpeckers. Exemplar species (i-iv) illustrate extreme variation in PC1: (i) exhibits high energy scores across most pattern element sizes, with small, medium, and large pattern elements; (ii) has low energy scores across the spectrum, with few pattern elements of any size. And, extreme variation in PC2: (iii) has many small pattern elements and few of any other sizes; (iv) has only medium and large size pattern elements. Illustrations © HBW Alive/Lynx Edicions.



Supplementary Fig. 18 | Principal component loadings of belly pattern. Principal component loadings of pattern energy spectra across filter size for all species reveals how belly pattern elements of different sizes influence PC scores.