

Supporting Information

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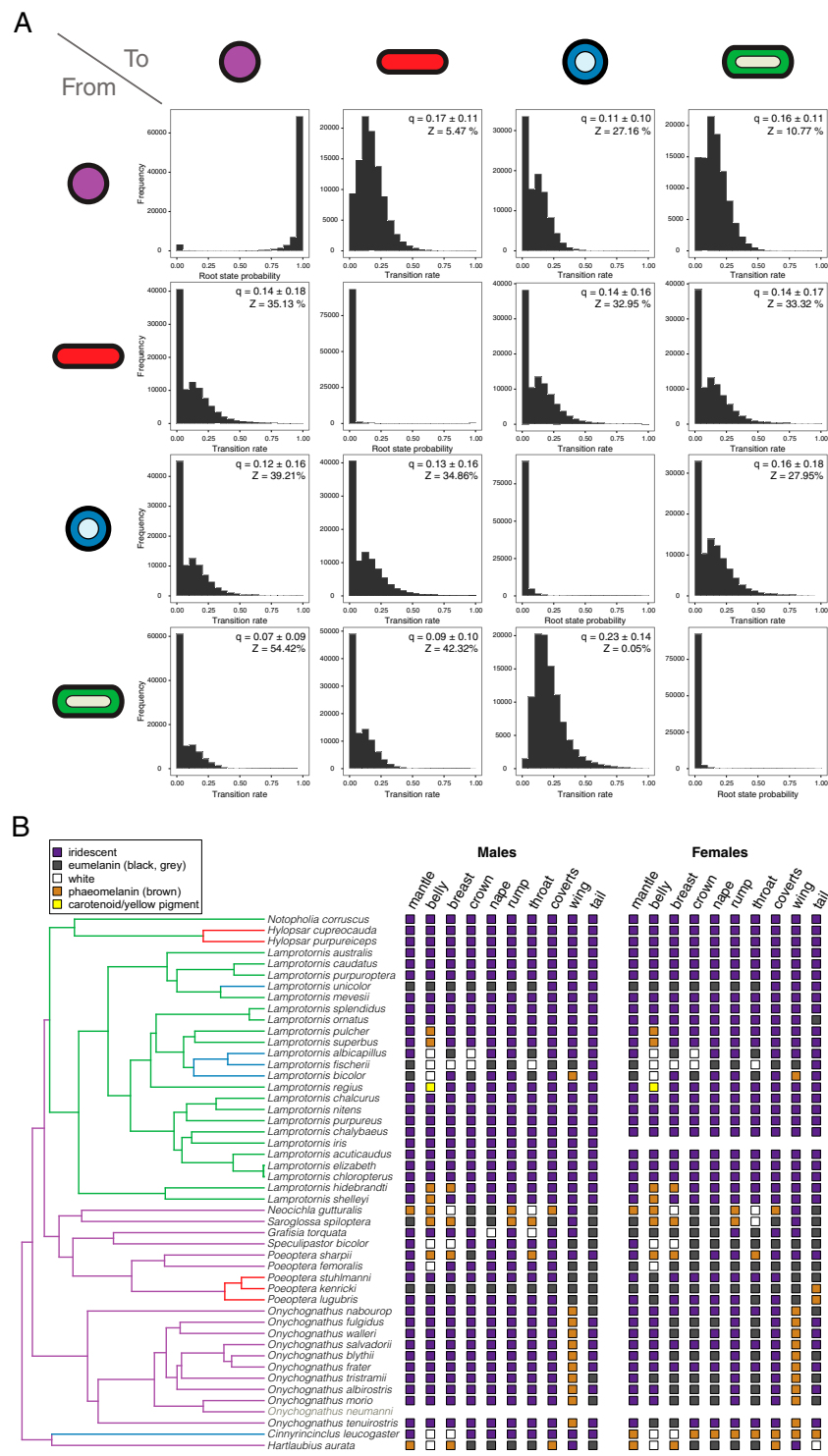


Fig. S1. (A) Posterior probabilities for the transition rates (off-diagonal) and root ancestral states (diagonal) for the different melanosome morphologies obtained from the reversible-jump Markov chain Monte Carlo analysis. Mean and SD for transition rates (q), as well as the frequency that the parameter was estimated to be zero (Z), are shown for each transition. (B) Maximum clade credibility tree for the African starlings, showing the melanosome regime used in evolutionary models for all color patches (Table S2) and the color phenotype observed for each sex of each species. *Onychognathus neumanni* and females of *Lamprolaima iris* were not sampled and therefore are not included in analyses.

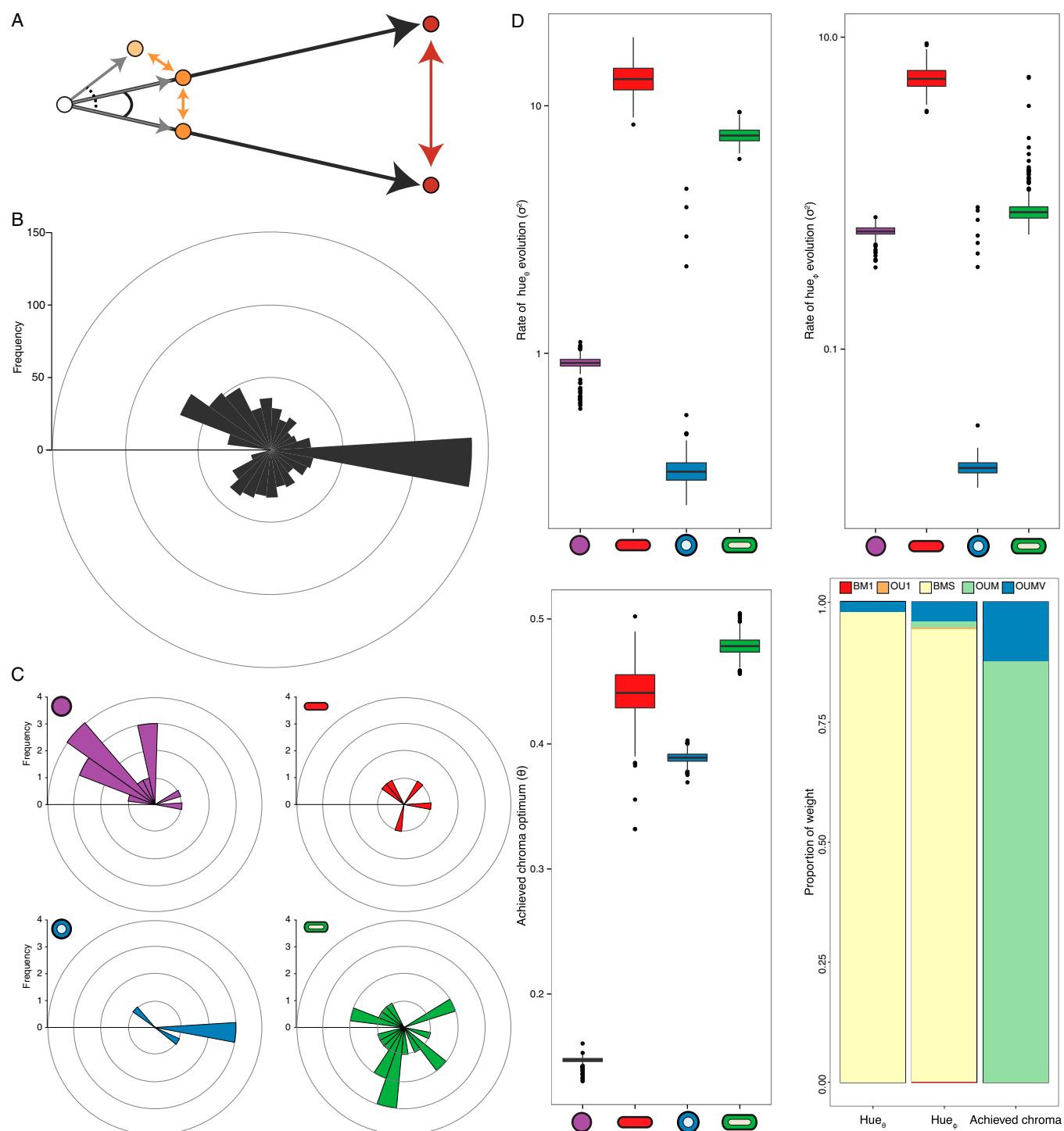


Fig. S2. Problems in using univariate descriptors of hue and saturation in describing color diversification. (A) In the tetrachromatic colorspace model, hue is described as the angles between the color point, the achromatic center (white circle), and either the *sml* cones plane (Hue_{e}) or the *u* vertex (Hue_{u}) and saturation as the distance between the color point and the achromatic center (black and gray arrows). If we define phenotypic disparity as the distance between points in colorspace, for a given hue difference (angle solid line), two color points can be very similar if close to the achromatic center (orange circles) or very different if these colors have higher chroma (i.e., distant from the achromatic center, red circles). Similarly, points with considerably different hue distances (dashed and solid angle lines) can be equidistant from each other (i.e., similar level of morphological disparity) depending on their distance from the achromatic center (as shown in the distance between the two dark orange circles and the light orange circles). (B) Further, as hue is described by angular variables, treating it as a linear variable results in an overestimation of the hue differences between points found near the zero angle (black line), which are treated as being at the extremes of a linear variable circular histogram of Hue_{e} angles for the iridescent colors across all body patches of African starlings, for both sexes. If measured colors spanned a restricted range of angles, treating this variable as a linear variable would be less problematic, but colors observed range across nearly all possible angles. (C) Circular histograms of Hue_{e} for the upper back of male African starlings by melanosome morphology, showing that this broad hue span is present even when a single body patch is considered. Therefore, the interactive effects of independent color descriptors were considered by summarizing color disparity in terms of multivariate orthogonal principal components for our analyses. (D) Parameter estimates and model selection results for the univariate descriptors hue and saturation for the male mantle patch color evolution across the 500 trees sampled from the posterior distribution.

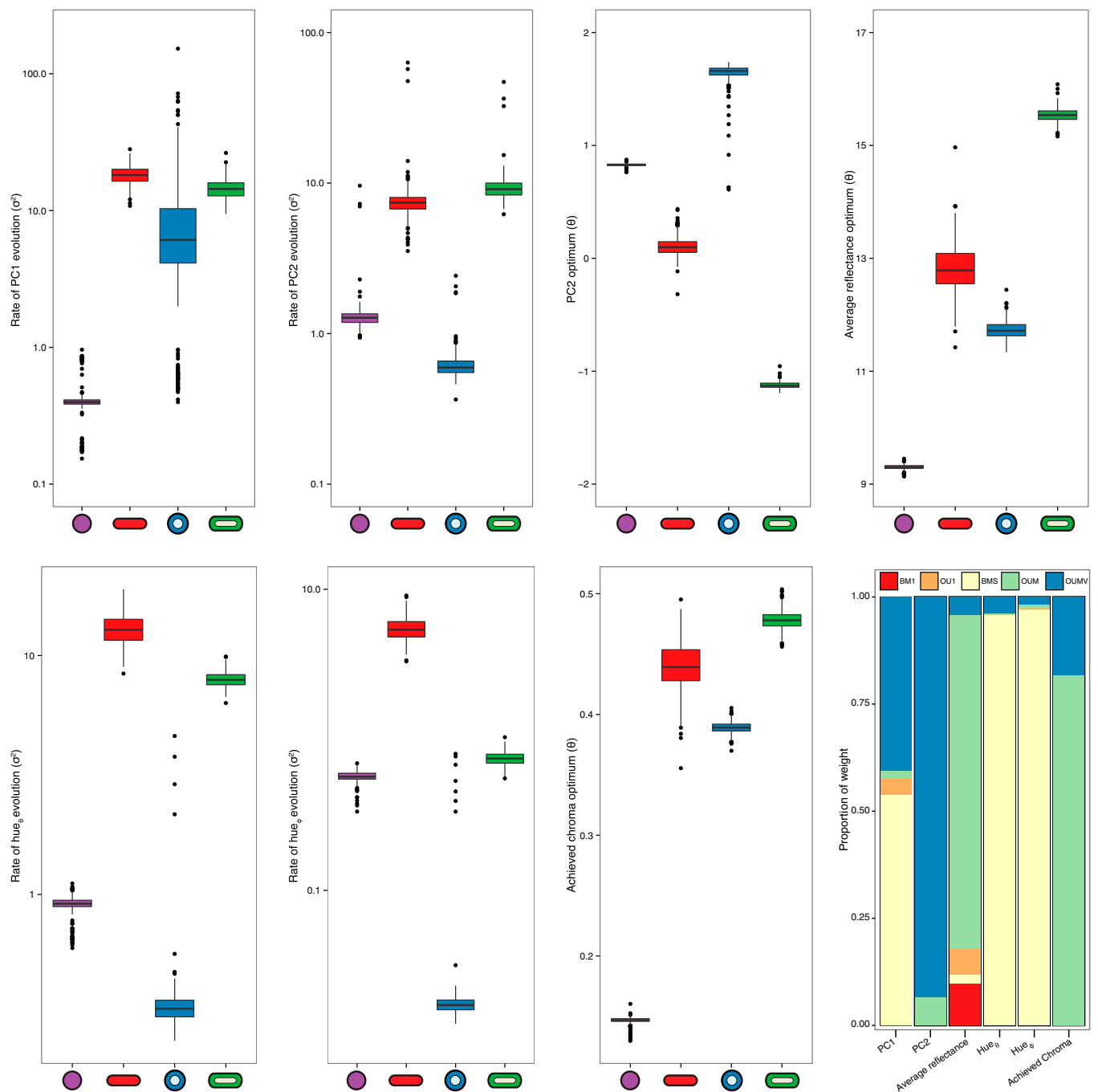


Fig. S3. Parameter estimates and model selection results under the best models for the principal components and univariate color descriptors for the male mantle patch color evolution across the 500 trees sampled from the posterior distribution, excluding *Lamprotornis elisabeth*.

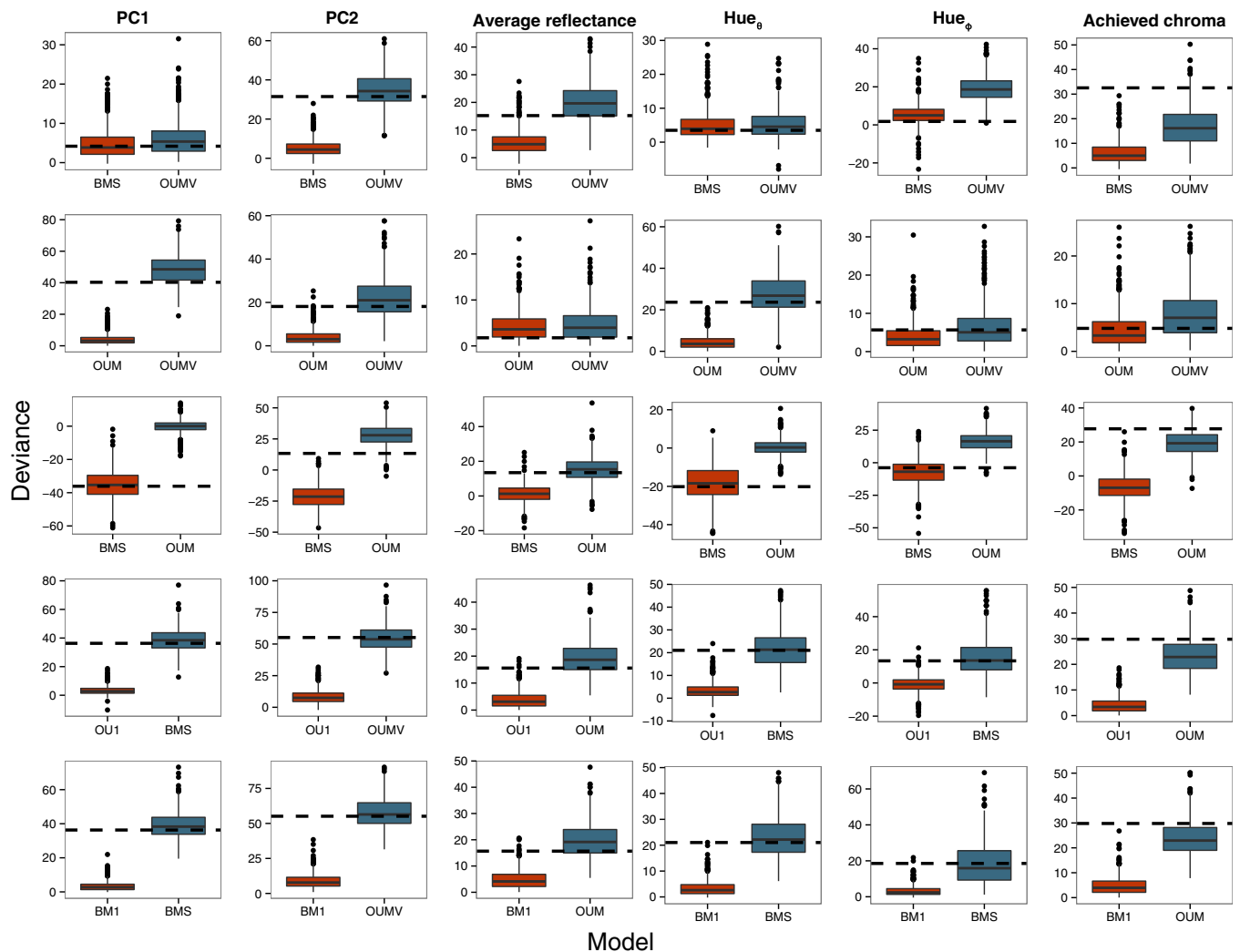


Fig. S5. Phylogenetic Monte Carlo results for the pairwise comparisons of evolutionary models for the male mantle patch variables. Boxes indicate the deviance value calculated by comparing the two models in the comparison after data were simulated under the respective model given the estimated maximum-likelihood estimates obtained from the maximum clade credibility tree (Fig. S1B). The dashed line indicates the deviance calculated from the maximum-likelihood models. Values below the line indicate low support for that model, and values that fall within the distribution obtained from a simulated model indicate a high probability of obtaining such deviance values under data simulated under that model. If the line falls within the distribution of both models, there is low power to distinguish between them, resulting from parameter estimates for different melanosome morphologies being similar for the parameters that distinguish both models (Table S2).

Table S1. Principal component analysis loadings and the variance explained by each component, obtained from the four cone stimuli that characterize avian color vision

	Component loadings			
Variable	PC1	PC2	PC3	PC4
Qu	0.52	0.46	−0.47	−0.54
Qs	0.46	−0.58	0.48	−0.48
Qm	−0.48	−0.52	−0.57	−0.41
Ql	−0.53	0.43	0.47	−0.56
Variance, %	64.43	31.22	04.35	<0.01

The first two components were used in the analyses. PC, principal component.

Table S3. Parameter estimates for the maximum clade credibility tree for all color patches and both sexes, as determined from AICc comparison (Table S2)

Table S4. Results from the maximum-likelihood multistate speciation and extinction (MuSSE) models under different combinations of parameters

Melanosome model	Transitions	Speciation rates	Speciation through time	log(Lik)	k	AICc	ΔAICc	λ ₁	λ ₂	λ ₃	λ ₄	λ _{slope}	q _{t12}	q _{t13}	q _{t14}	q _{t21}	q _{t23}	q _{t24}	q _{t31}	q _{t32}	q _{t34}	q _{a11}	q _{a2}	q _{a3}	
													q _{t12}	q _{t13}	q _{t14}	q _{t21}	q _{t23}	q _{t24}	q _{t31}	q _{t32}	q _{t34}	q _{a11}	q _{a2}	q _{a3}	
Irreversible Full	Single	Derived	Linear	-31.58	4	72.12	0	-0.24	1.11	1.11	1.11	5.44	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13	
	Single	Derived	Linear	-32.65	4	74.26	2.14	-0.24	1.11	1.11	1.11	5.43	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	
	Asymmetric	Derived	Linear	-31.52	5	74.51	2.39	-0.24	1.1	1.1	1.1	5.42	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
Irreversible	Single	Single	Linear	-34.69	3	75.94	3.82	0.64	0.64	0.64	0.64	4.69	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13	
	Single	Full	Linear	-31.44	6	76.99	4.87	-0.24	1.3	0.45	1.16	5.45	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13	
	Asymmetric	Derived	Linear	-31.52	6	77.15	5.03	-0.24	1.1	1.1	1.1	5.42	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
Stepwise Full	Single	Derived	Linear	-34.21	4	77.37	5.25	-0.22	1.07	1.07	1.07	5.38	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19	
	Single	Single	Linear	-35.76	3	78.07	5.95	0.64	0.64	0.64	0.64	4.69	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	
	Asymmetric	Single	Linear	-34.59	4	78.12	6	0.64	0.64	0.64	0.64	4.69	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
Irreversible Full	Single	Full	Linear	-32.52	6	79.13	7.01	-0.24	1.31	0.46	1.16	5.44	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	
	Asymmetric	Full	Linear	-31.39	7	79.65	7.53	-0.24	1.3	0.45	1.16	5.43	0.12	0.12	0.12	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
	Asymmetric	Single	Linear	-34.59	5	80.63	8.51	0.64	0.64	0.64	0.64	4.69	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
Stepwise	Single	Single	Linear	-37.11	3	80.78	8.66	0.64	0.64	0.64	0.64	4.69	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19	
	Single	Full	Linear	-34.02	6	82.13	10.01	-0.21	0.84	0.42	1.24	5.37	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19	
	Asymmetric	Full	Linear	-31.39	8	82.56	10.44	-0.24	1.3	0.45	1.16	5.43	0.12	0.12	0.12	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
Irreversible	Single	Single	Constant	-40.5	2	85.28	13.16	2.1	2.1	2.1	2.1	0	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13	
	Single	Derived	Constant	-39.9	3	86.35	14.23	1.76	2.46	2.46	2.46	0	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13	
	Asymmetric	Single	Constant	-40.4	3	87.35	15.23	2.1	2.1	2.1	2.1	0	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
Irreversible Full	Single	Single	Constant	-41.57	2	87.41	15.29	2.1	2.1	2.1	2.1	0	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	
	Single	Derived	Constant	-40.96	3	88.48	16.36	1.76	2.46	2.46	2.46	0	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	
	Asymmetric	Derived	Constant	-39.78	4	88.51	16.39	1.76	2.46	2.46	2.46	0	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
Stepwise	Full	Derived	Linear	-29.88	11	89.31	17.19	-0.28	1.17	1.17	1.17	5.51	0.17	0.26	0	1.0	0	0	0	0	3.34	0	0	0.48	
	Asymmetric	Single	Constant	-40.4	4	89.74	17.62	2.1	2.1	2.1	2.1	0	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
	Single	Single	Constant	-42.92	2	90.11	17.99	2.1	2.1	2.1	2.1	0	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19	
Irreversible	Single	Full	Constant	-39.54	5	90.53	18.41	1.77	2.13	1.45	2.66	0	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13	
	Single	Single	Linear	-32.38	10	90.87	18.75	0.64	0.64	0.64	0.64	4.69	0	0.06	0	1.55	0	0.57	0	0	0	0	0	0.36	
	Asymmetric	Derived	Constant	-39.78	5	91.02	18.9	1.76	2.46	2.46	2.46	0	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16	
Stepwise Full	Single	Derived	Constant	-40.61	3	91.19	19.07	1.76	2.45	2.45	2.45	0	0.18	0.18	0	0.18	0	0.18	0.18	0	0.18	0	0.18	0.18	
	Single	Full	Constant	-42.31	5	92.67	20.55	1.78	2.13	1.46	2.66	0	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	
	Asymmetric	Derived	Linear	-29.88	12	92.93	20.81	-0.27	1.15	1.15	1.15	5.49	0.15	0.25	0	0	0	0	0	0.08	3.22	0	0	0.48	
Irreversible	Full	Derived	Constant	-39.42	6	92.93	20.81	1.77	2.12	1.44	2.66	0	0.11	0.11	0.11	0	0.17	0.17	0	0.17	0.17	0	0.17	0.17	
	Asymmetric	Full	Constant	-39.42	6	92.93	20.81	1.77	2.12	1.44	2.66	0	0.11	0.11	0.11	0	0.17	0.17	0	0.17	0.17	0	0.17	0.17	
	Single	Full	Constant	-41.89	5	95.24	23.12	1.77	2.19	1.38	2.69	0	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19	
Stepwise	Single	Single	Linear	-32.94	11	95.43	23.31	0.64	0.64	0.64	0.64	4.69	0.21	0.08	0	0	0	1.14	0	0	0	0	0	0.31	
	Single	Full	Constant	-39.42	7	95.7	23.58	1.77	2.12	1.44	2.66	0	0.11	0.11	0.11	0	0.17	0.17	0	0.17	0.17	0	0.17	0.17	
	Asymmetric	Full	Constant	-39.42	7	95.7	23.58	1.77	2.12	1.44	2.66	0	0.11	0.11	0.11	0	0.17	0.17	0	0.17	0.17	0	0.17	0.17	
Irreversible Full	Full	Full	Linear	-29.81	13	96.65	24.53	-0.21	0.79	0.4	1.26	5.38	0.21	0.09	0	0	0	1.15	0	0	0	0	0	0.32	
	Full	Derived	Constant	-36.06	10	98.23	26.11	1.28	2.63	2.63	2.63	0	0	0.08	0	1.73	0	0.53	0	0	0	0	0	0.33	
	Full	Single	Constant	-38.19	9	99.24	27.12	2.1	2.1	2.1	2.1	0	0	0.06	0	1.55	0	0.57	0	0	0	0	0	0.36	
Stepwise	Full	Single	Linear	-29.81	14	100.74	28.62	-0.21	0.79	0.4	1.26	5.38	0.21	0.09	0	0	0	1.15	0	0	0	0	0	0.32	
	Full	Single	Constant	-38.75	10	103.62	31.5	2.1	2.1	2.1	2.1	0	0.21	0.08	0	0	0	1.14	0	0	0	0	0	0.31	
	Full	Full	Constant	-35.56	12	104.3	32.18	0	3.76	3.86	1.78	0	0	0.43	0	3.29	0	0	0	0	3.42	0	0	0.09	
Irreversible	Full	Derived	Linear	-29.88	15	105.24	33.12	-0.27	1.15	1.15	1.15	5.49	0.15	0.25	0	0	0	1.14	0	0	0.08	3.22	0	0	0.48
	Full	Single	Constant	-38.11	11	105.77	33.65	1.75	2.46	2.46	2.46	0	0.2	0.09	0	0	0	1.14	0	0	0	0	0	0.35	
	Full	Single	Constant	-32.34	14	105.8	33.68	0.64	0.64	0.64	0.64	4.69	0	0	1.43	0.11	0.54	0	0	0	0	0	0	0.35	
Irreversible Full	Full	Full	Constant	-37.56	13	112.16	40.04	1.72	2.24	4.03	1.51	0	0.1	0.2	0	0	0	0	0	0	3.32	0	0.16	0	

Table S4. Cont.

Melanosome model	Transitions	Speciation rates	Speciation through time	log(Lik)	k	AICc	ΔAICc	λ ₁	λ ₂	λ ₃	λ ₄	λ _{slope}	q _{t12}	q _{t13}	q _{t14}	q _{t21}	q _{t23}	q _{t24}	q _{t31}	q _{t32}	q _{t34}	q _{t41}	q _{t42}	q _{t43}
Full	Full	Derived	Constant	-36.06	14	113.24	41.12	1.28	2.63	2.63	2.63	0	0	0.08	0	1.73	0	0.53	0	0	0	0	0	0.33
Full	Full	Single	Constant	-38.15	13	113.32	41.2	2.1	2.1	2.1	2.1	0	0	0	0	1.43	0.11	0.54	0	0	0	0	0	0.35
Full	Full	Full	Linear	-29.81	17	114.72	42.6	-0.21	0.79	0.4	1.26	5.38	0.21	0.09	0	0	0	1.15	0	0	0	0	0	0.32
Full	Full	Full	Constant	-35.37	16	120.88	48.76	0	3.76	3.83	1.77	0	0	0.33	0	3.17	0.12	0	0	0	3.39	0	0	0.09
Excluding <i>Lamprolornis elisabeth</i>																								
Irreversible	Single	Derived	Linear	-31.55	4	72.06	0	-0.3	0.82	0.82	0.82	5.82	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13
Full	Single	Derived	Linear	-32.63	4	74.2	2.14	-0.32	0.81	0.81	0.81	5.85	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11
Irreversible	Asymmetric	Derived	Linear	-31.49	5	74.44	2.38	-0.31	0.81	0.81	0.81	5.84	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Irreversible	Single	Single	Linear	-33.98	3	74.52	2.46	0.31	0.31	0.31	0.31	5.62	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13
Full	Single	Single	Linear	-35.05	3	76.65	4.59	0.31	0.31	0.31	0.31	5.62	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11
Irreversible	Asymmetric	Single	Linear	-33.88	4	76.71	4.65	0.31	0.31	0.31	0.31	5.62	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Irreversible	Single	Full	Linear	-31.43	6	76.96	4.9	-0.33	1.25	0.37	0.78	5.9	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13
Full	Asymmetric	Derived	Linear	-31.49	6	77.08	5.02	-0.31	0.81	0.81	0.81	5.84	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Stepwise	Single	Derived	Linear	-34.15	4	77.25	5.19	-0.3	0.78	0.78	0.78	5.81	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19
Full	Single	Full	Linear	-32.5	6	79.1	7.04	-0.32	1.25	0.38	0.77	5.89	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11
Full	Asymmetric	Single	Linear	-33.88	5	79.22	7.16	0.31	0.31	0.31	0.31	5.62	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Stepwise	Single	Single	Linear	-36.4	3	79.36	7.3	0.31	0.31	0.31	0.31	5.62	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19
Irreversible	Asymmetric	Full	Linear	-31.36	7	79.6	7.54	-0.31	1.24	0.38	0.78	5.85	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Stepwise	Single	Full	Linear	-34.08	6	82.26	10.2	-0.29	0.78	0.35	0.86	5.79	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19
Full	Asymmetric	Full	Linear	-31.36	8	82.52	10.46	-0.31	1.24	0.38	0.78	5.85	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Irreversible	Single	Single	Constant	-41.23	2	86.73	14.67	2.06	2.06	2.06	2.06	0	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13
Irreversible	Single	Derived	Constant	-40.77	3	88.1	16.04	1.76	2.36	2.36	2.36	0	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13
Irreversible	Asymmetric	Single	Constant	-41.12	3	88.81	16.75	2.06	2.06	2.06	2.06	0	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Full	Single	Single	Constant	-42.29	2	88.86	16.8	2.06	2.06	2.06	2.06	0	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11
Stepwise	Full	Derived	Linear	-29.91	11	89.37	17.31	-0.35	0.87	0.87	0.87	5.91	0.17	0.26	0	0	0	0	0	0	3.32	0	0	0.48
Stepwise	Full	Single	Linear	-31.67	10	89.45	17.39	0.31	0.31	0.31	0.31	5.62	0	0.06	0	1.55	0	0.57	0	0	0	0	0	0.36
Full	Single	Derived	Constant	-41.84	3	90.23	18.17	1.76	2.36	2.36	2.36	0	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11
Irreversible	Asymmetric	Derived	Constant	-40.65	4	90.26	18.2	1.76	2.37	2.37	2.37	0	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Full	Asymmetric	Single	Constant	-41.12	4	91.2	19.14	2.06	2.06	2.06	2.06	0	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Stepwise	Single	Single	Constant	-43.65	2	91.57	19.51	2.06	2.06	2.06	2.06	0	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19
Irreversible	Single	Full	Constant	-40.49	5	92.44	20.38	1.77	2.13	1.46	2.54	0	0.13	0.13	0.13	0	0.13	0.13	0	0.13	0.13	0	0.13	0.13
Full	Asymmetric	Derived	Constant	-40.65	5	92.77	20.71	1.76	2.37	2.37	2.37	0	0.11	0.11	0.11	0	0.16	0.16	0	0.16	0.16	0	0.16	0.16
Stepwise	Single	Derived	Constant	-43.19	3	92.93	20.87	1.76	2.36	2.36	2.36	0	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19
Irreversible	Full	Derived	Linear	-29.89	12	92.95	20.89	-0.34	0.85	0.85	0.85	5.87	0.14	0.24	0	0	0	0	0	0.15	3.11	0	0	0.49
Irreversible	Full	Single	Linear	-32.23	11	94.01	21.95	0.31	0.31	0.31	0.31	5.62	0.21	0.08	0	0	0	1.14	0	0	0	0	0	0.31
Full	Single	Full	Constant	-41.55	5	94.57	22.51	1.78	2.13	1.46	2.53	0	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11	0.11
Irreversible	Asymmetric	Full	Constant	-40.37	6	94.83	22.77	1.77	2.13	1.45	2.54	0	0.11	0.11	0.11	0	0.17	0.17	0	0.17	0.17	0	0.17	0.17
Stepwise	Full	Full	Linear	-29.8	13	96.63	24.57	-0.35	1.32	1.11	0.69	5.89	0.17	0.26	0	0	0	0	0	0	3.35	0	0	0.43
Stepwise	Single	Full	Constant	-42.85	5	97.16	25.1	1.77	2.2	1.38	2.55	0	0.19	0.19	0	0.19	0	0.19	0.19	0	0.19	0	0.19	0.19
Full	Asymmetric	Full	Constant	-40.37	7	97.6	25.54	1.77	2.13	1.45	2.54	0	0.11	0.11	0.11	0	0.17	0.17	0	0.17	0.17	0	0.17	0.17
Stepwise	Full	Derived	Constant	-37.01	10	100.13	28.07	1.28	2.55	2.55	2.55	0	0	0.08	0	1.72	0	0.53	0	0	0	0	0	0.33
Stepwise	Full	Single	Constant	-38.92	9	100.7	28.64	2.06	2.06	2.06	2.06	0	0	0.06	0	1.55	0	0.57	0	0	0	0	0	0.36
Irreversible	Full	Full	Linear	-29.8	14	100.72	28.66	-0.35	1.3	0.99	0.7	5.9	0.15	0.25	0	0	0	0	0	0.09	3.23	0	0	0.45
Full	Full	Single	Linear	-31.63	14	104.38	32.32	0.31	0.31	0.31	0.31	5.62	0	0	0	1.43	0.11	0.54	0	0	0	0	0	0.33

Speciation
through

Estimates represent the maximum-likelihood estimates obtained under each model after 5,000 optimization procedures starting at different points. Melanosome models considered were full (all transitions estimated), irreversible (transitions back to the ancestral state set to zero), and stepwise (transitions involving both a change in shape and hollowiness set to zero); transition models considered were full (all rates different), asymmetric (two transition rates, with transitions between derived morphologies different from transitions to/from ancestral morphology), and single (all transition rates equal); speciation rate models considered were single (all melanosome morphologies share the same speciation rate), derived (derived melanosomes share the same speciation rate, which is different from the ancestral melanosome speciation rate), and full (rates different for all four melanosome morphologies); and speciation through time models were constant and linear over time. Results are shown for all species and excluding *Lamprolornis elisabeth*.