

# Carotenoid-Based Ornaments of Female and Male American Goldfinches (*Spinus tristis*) Show Sex-Specific Correlations with Immune Function and Metabolic Rate

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## ABSTRACT

Conspicuous ornamentation has been linked to immunological and physiological condition in males of many species. In species where both sexes are ornamented, it is unclear whether the signal content of ornaments differs between males and females. We examined the immunological and physiological correlates of carotenoid-based bill and plumage ornamentation in American goldfinches *Spinus tristis*, a species in which bright orange bills are sexually monomorphic but yellow plumage is sexually dimorphic during the breeding season. Because bill color is dynamic over short periods while plumage color is static over longer time frames, we tested whether these signals have the potential to provide temporal information about immunity and condition. In both sexes, bill color (but not plumage color) was negatively related to leukocyte differential, a measure of recent stress, while plumage color (but not bill color) was positively related to resting metabolic rate. In females, bill color also positively correlated with immunoglobulin Y, a component of acquired immunity, while plumage color positively predicted natural antibody levels, a component of innate immunity. In males, neither bill color nor plumage color predicted immune function, suggesting that the mechanisms underlying these signals vary with sex. Our results demonstrate that dynamic signals such as bill coloration do not merely reflect the same information provided by static signals but that these two classes of

signal provide information about different temporal aspects of phenotypic quality. Furthermore, our results indicate that a signal expressed in both sexes has the potential to provide different information depending on the sex of the bearer.

## Introduction

In many animals, males display conspicuous ornamentation, and the most elaborate males achieve the highest mating success (Andersson 1994). Extensive evidence supports a link between male signal strength and indexes of immunity across a wide variety of taxa, including birds (e.g., Dufva and Allander 1995; Doucet and Montgomerie 2003; Faivre et al. 2003; McGraw and Ardia 2003; Dunn et al. 2010), fish (Clotfelter et al. 2007), and lizards (Martín and López 2009). There is increasing evidence that elaborate female traits also confer a selective advantage during inter- or intrasexual interactions (see Amundsen 2000; LeBas 2006; Clutton-Brock 2009). Female ornaments have been found to signal immune function (but see Pärn et al. 2005), yet most studies on females have focused on species in which ornamental expression is limited to or more pronounced in females (Roulin et al. 2001a, 2001b; Hanssen et al. 2006). In contrast, few studies have examined whether ornaments shared by the sexes signal the same or different aspects of immunity or physiological condition (but see Alvarez et al. 2005; Lopez et al. 2008; Maney et al. 2008). The observation that males and females of a species often differ in immunocompetence (e.g., Nunn et al. 2009; Pap et al. 2010) raises the question of whether ornamentation shared by the sexes reflects immune function in the same way in both sexes. Given that ornaments may be maintained by different selective forces acting on each sex (Heinsohn et al. 2005; LeBas 2006; Murphy 2006, 2007; Murphy et al. 2009), we might expect the information signaled by ornaments to differ between males and females.

The vertebrate immune system can be broadly divided into innate and acquired (or adaptive) immunity components (e.g., Demas et al. 2011). The innate immune system provides the first line of defense against infection and contributes to rapid nonspecific protection against invading pathogens through both humoral and cellular components. The acquired immune system is divided into humoral and cell-mediated immunity and initiates slower responses that target specific infectious agents, providing long-lasting protection against bacteria, vi-

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ruses, fungi, and other parasites. Increasingly, researchers recognize that in order to gain a more complete understanding of an individual animal's immunocompetence, it is necessary to assess components of innate and acquired arms simultaneously, because allocation to one arm of the immune system may or may not be correlated with allocation to the other (Norris and Evans 2000; Saks et al. 2003; Hörak et al. 2006; Buehler et al. 2008). Thus, investment in innate versus acquired immunity may vary with sex, age, or investment in other life-history traits (Martin et al. 2007; Ardia et al. 2011). As such, the two different arms may provide contrasting views of the relationship between ornamentation and immune function, and in cases where an ornament is shared by the sexes, it seems plausible that the ornament may signal different components of immunity in males and females.

Carotenoid-based ornamentation often signals immunocompetence (e.g., Saks et al. 2003; Aguilera and Amat 2007; Baeta et al. 2008; Dunn et al. 2010). Increased intake of dietary carotenoids has been associated with various immune function benefits, including increased cell-mediated and humoral immune responsiveness and decreased DNA damage and inflammation (e.g., Chew et al. 2011). However, the specific effects of carotenoids on the immune response vary with type of carotenoid, dosage, and species studied (Chew and Park 2004). Links between carotenoids and health benefits are likely due to their antioxidant properties, although other mechanisms, including gap junction communication and regulation of membrane fluidity, are also likely involved (Chew and Park 2004; Rao and Rao 2007). Carotenoid-based ornaments may thus allow an individual to signal health because allocation of carotenoids to ornamentation can occur only if carotenoids are not being used elsewhere (e.g., Lozano 1994; Møller et al. 2000; but see Costantini and Møller 2008). Alternatively, carotenoid ornamentation may be correlated with immune function because both are dependent on the functionality of a common set of biochemical pathways that are themselves dependent on individual condition (reviewed by Hill 2011). Regardless of the underlying mechanism, these hypotheses predict that carotenoid ornamentation should reliably signal immunocompetence.

Immune function has also been linked with mass-adjusted basal (or resting) metabolic rate (Tieleman et al. 2005), although this relationship is complex (see Martin et al. 2007; Tieleman et al. 2008). Among species, variation in resting metabolic rate (RMR) reflects the "pace of life" such that species that tend to invest in survival over current reproduction tend to have lower RMR (Tieleman et al. 2005; Wiersma et al. 2007). However, among individuals within a species, RMR may indicate an individual's capacity to perform well during challenging activities (e.g., Biro and Stamps 2010). For example, some components of immune function, such as constitutive innate immunity, have been found to positively relate to mass-adjusted RMR within a species (Tieleman et al. 2005). Additionally, experimental activation of the immune system can lead to an elevation in RMR, presumably to fuel the costs of the immune activity (e.g., Burness et al. 2010; Van de Crommen-

acker et al. 2010). Individuals in higher condition should be able to expend more energy toward both immunity and ornamentation, so we expect high-quality individuals to be able to sustain both high RMR and high ornamentation.

In this study we examined the immunological and metabolic correlates of carotenoid-based bill and plumage ornamentation in male and female American goldfinches *Spinus tristis*. In this species, both males and females express a common set of ornaments, but their ornaments appear to function as communication signals in different social contexts. Orange bill coloration, which is expressed similarly in the sexes, functions in females as a signal of status during competitive interactions with other females (Murphy et al. 2009), while male bill color does not appear to play a signaling role in the same competitive context (T. G. Murphy and K. A. Tarvin, unpublished data). Moreover, experimental manipulations of bill color indicate that it does not play a role in mate choice in either sex (T. G. Murphy and K. A. Tarvin, unpublished data). A second conspicuous ornament is the saturated yellow plumage, which is more pronounced in males. Male plumage color functions as a mate-choice signal: in an aviary mate-choice trial, Johnson et al. (1993) found that females preferred males with darker throat plumage (i.e., lower luminance). In contrast, the function of female plumage color remains unknown. However, based on findings of assortative mating with respect to plumage color saturation (MacDougall and Montgomerie 2003), female plumage may also function as a mate-choice signal. In addition to playing different roles in communication, these two ornaments vary in an important way: bill coloration has been found to change rapidly, in a matter of hours, under experimentally induced physiological perturbations in both sexes (Rosen and Tarvin 2006; M. F. Rosenthal, T. G. Murphy, N. Darling, and K. A. Tarvin, unpublished data), whereas plumage color changes little between the molt in early spring, when they acquire breeding plumage, and the loss of their colorful feathers after the breeding season. Thus, dynamic bill color has the potential to reflect current physiological state, whereas static plumage color can reflect only condition at the time of molt.

We hypothesize that bill and plumage color are condition-dependent signals in both sexes and that dynamic bill color reflects aspects of condition that fluctuate over a short time frame whereas static plumage color reflects less labile components of condition. Furthermore, based on the different roles these ornaments appear to play in male and female communication (Johnson et al. 1993; Murphy et al. 2009), we explore the possibility that these shared ornaments communicate different aspects of immunity and condition in males and females. If the ornaments do indeed have sex-specific roles in signaling, it seems possible that differences exist between the sexes in what the ornaments reflect; however, it is also possible that the ornaments reflect the same component of condition in both sexes and that receivers respond to the ornaments in a sex-specific manner (e.g., disregarding information conveyed in a signal if given by one sex while assessing it in the other sex). Because both are plausible explanations, we make no directional predictions about sex differences and simply predict that highly

ornamented goldfinches (of both sexes) will have lower stress, higher RMR, and more robust innate and adaptive immune systems. We measured stress based on the ratio of heterophils to lymphocytes (H:L) in whole blood because high ratios of these immunologically important leukocytes are indicative of stress and have been positively correlated with susceptibility to infection and negatively correlated with growth rate and survival (reviewed in Davis et al. 2008; see also Müller et al. 2011). We additionally test the prediction that ornamental coloration is positively correlated with RMR because individuals with higher RMR are expected to have a greater capacity to engage in energetically costly activities (e.g., parental care, competitive interactions). To assess immunity, we measured levels of natural antibodies (NAb) and immunoglobulins (IgY). NAb, a component of the innate humoral immune system, bind to foreign antigens and neutralize infection (Ochsenbein and Zinkernagel 2000). IgY, a component of the adaptive humoral immune system, is the most abundant class of antibodies in avian blood and on infection coats microorganisms, allowing other immune cells to recognize and destroy them (Demas et al. 2011).

## Material and Methods

### *Species*

Both male and female American goldfinches develop vibrant orange bills and moult into colorful yellow plumage before the onset of the breeding season. Although bill color is very similar in the two sexes, their plumage is markedly sexually dichromatic: the yellow plumage of males covers much of the body whereas that of females tends to be drabber and occurs in small patches, though the size of the patches is highly variable (Rosen and Tarvin 2006; McGraw and Middleton 2009).

In previous research, we found that bill color of wild male and female goldfinches held in captivity significantly declined within 7 h of capture (presumably because of stress of captivity). The decline continued over several days and was significantly greater among birds (males and females) that were exposed to an experimental immune challenge (M. F. Rosenthal, T. G. Murphy, N. Darling, and K. A. Tarvin, unpublished data). Thus, bill color is dynamic over short time periods, and fluctuations in bill color appear to correspond with short-term changes in physiological condition.

### *General Methods*

We captured 41 male and 28 female American goldfinches in July 2008 at Queens University Biological Station in Ontario, Canada (44°33'N, 76°19'W). Goldfinches were captured with mist nets placed around feeders at three different sites within a 15-km radius. On capture, we categorized sex and age class (in their second calendar year [SY]—i.e., hatched the previous summer—and after their second year [ASY]) based on plumage (Pyle 1997), measured mass ( $\pm 0.1$  g), and morphological features ( $\pm 0.1$  mm) and collected ca. 100  $\mu$ L of blood. Blood was collected from the brachial wing vein into two heparinized capillary tubes, sealed with clay, and stored up to 4 h on ice

until processing. An additional drop of blood was collected into a nonheparinized capillary tube and used to make two blood smears following Davis (2005). Slides were air-dried and stored until staining took place. Blood samples were centrifuged (Hemata STAT-II, Separation Technology) for three 60-s intervals. Plasma was removed, placed into microcentrifuge tubes, and frozen at  $-20^{\circ}\text{C}$  for up to 4 wk. Samples were then transferred to a  $-80^{\circ}\text{C}$  freezer until analysis.

Because bill color is dynamic and changes rapidly in this species in response to stress of capture (Rosen and Tarvin 2006; M. F. Rosenthal, T. G. Murphy, N. Darling, and K. A. Tarvin, unpublished data), we measured color reflectance ( $R$ ) of the bill within 1 h of capture (mean  $\pm$  SE =  $32.1 \pm 2.7$  min), which is a few hours before stress-induced color change has been detected spectrometrically (M. F. Rosenthal, T. G. Murphy, N. Darling, and K. A. Tarvin, unpublished data). Individuals were then transported to the biological station and placed in visually isolated housing units (1.2 m<sup>3</sup>) for ca. 6 h, where they received water and Nyjer seed ad lib. On the same evening of capture, up to three birds were placed into individual metabolic chambers where oxygen consumption was measured overnight as an index of RMR. After RMR was measured, birds were used in a dominance experiment the next morning (see Murphy et al. 2009) and released at their capture site. Only adult (ASY) birds were used for this experiment. All research was conducted under Queen's University animal care protocol 2005-044 and Trent University protocol 08046.

### *Color Analysis*

We collected spectral measures from the bill and throat of each bird. We used an Ocean Optics USB2000+ spectrometer and a PX-2 pulsed xenon lamp (Ocean Optics, Dunedin, FL) with the probe held at  $90^{\circ}$  to the color patch. The probe was fixed within a dark plastic holder, which held the probe tip approximately 7 mm from the color patch. We quantified  $R$  as the proportion of light reflected from the color patch compared with a Spectralon white standard (Labsphere) at 1-nm intervals across the avian visual range (320–700 nm). We calculated the mean  $R$  of five measures for each color patch, each taken at a different haphazardly chosen location within a color patch. Using mean  $R$  curves, we calculated mean luminance ("brightness"; mean  $R$  from 320 to 700 nm); hue, a measure of spectral location (wavelength where  $R = [R_{\max} + R_{\min}]/2$ ); and yellow saturation, a measure of spectral purity (sum of  $R$  from 550 to 625 nm/mean luminance) using CLR 1.05 (Montgomerie 2008; see table 3.2 in Montgomerie 2006 for further details). These color analysis techniques are standard for carotenoid-based traits (Montgomerie 2008) and are similar to those used in other studies of goldfinches (MacDougall and Montgomerie 2003; Hill et al. 2009). All color measures were taken by T.G.M.

### *Natural Antibody Levels*

Innate humoral immunity was measured by quantifying NAb and complement in each individual's plasma following Matson

Table 1: Color and physiological parameters of male and female American goldfinches

Variable	Mean $\pm$ SE ( <i>n</i> )		Capture date		Sex	
	Males	Females	<i>F</i>	<i>P</i>	<i>F</i>	<i>P</i>
Bill luminance	.266 $\pm$ .007 (32)	.222 $\pm$ .008 (25)	.64	.426	16.79	<.001
Bill saturation	.248 $\pm$ .001 (32)	.246 $\pm$ .002 (25)	2.69	.107	1.81	.184
Bill hue	550.1 $\pm$ .957 (32)	546.2 $\pm$ 1.085 (25)	23.81	<.001	7.23	.010
Throat luminance	.302 $\pm$ .005 (32)	.260 $\pm$ .006 (25)	6.92	.011	31.18	<.001
Throat saturation	.324 $\pm$ .002 (32)	.300 $\pm$ .003 (25)	17.02	<.001	44.81	<.001
Throat hue	505.7 $\pm$ .454 (32)	501.0 $\pm$ .515 (25)	11.80	.001	46.42	<.001
IgY	.148 $\pm$ .007 (24)	.171 $\pm$ .007 (20)	1.28	.265	6.11	.018
NAb	1.191 $\pm$ .123 (19)	1.745 $\pm$ .126 (18)	7.75	.009	9.69	.004
H : L ratio	.768 $\pm$ .076 (28)	.579 $\pm$ .065 (22)	1.44	.236	2.44	.125
RMR <sup>a</sup>	43.910 $\pm$ .673 (24)	43.510 $\pm$ .793 (18)	.01	.926	.13	.719

Note. Estimated marginal means and standard errors are presented when the effect of capture date was significant. Unadjusted means and standard errors are presented otherwise. IgY = immunoglobulin Y; NAb = natural antibodies; H : L ratio = heterophil-to-lymphocyte ratio; RMR = resting metabolic rate.

<sup>a</sup>Estimated marginal means and standard error adjusted for effect of mass:  $F = 31.81$ ,  $P < 0.001$ .

et al. (2005). Because complement was detected for only a single individual, we do not discuss complement further. NAb titers were obtained for 37 adults (19 males, 18 females). Intra- and interplate assay variation for hemagglutination assays (obtained from two control samples run on each assay plate) was 3.3% and 2.5%, respectively. The assay and scoring were performed blindly by R.J.K.

#### Immunoglobulin Levels

Adaptive humoral immunity was estimated by assaying the concentration of immunoglobulin Y (IgY). We measured plasma levels of IgY using an enzyme-linked immunoabsorbent assay following Bourgeon et al. (2006) using goldfinch-specific plasma dilutions. All plasma samples were run in duplicate,

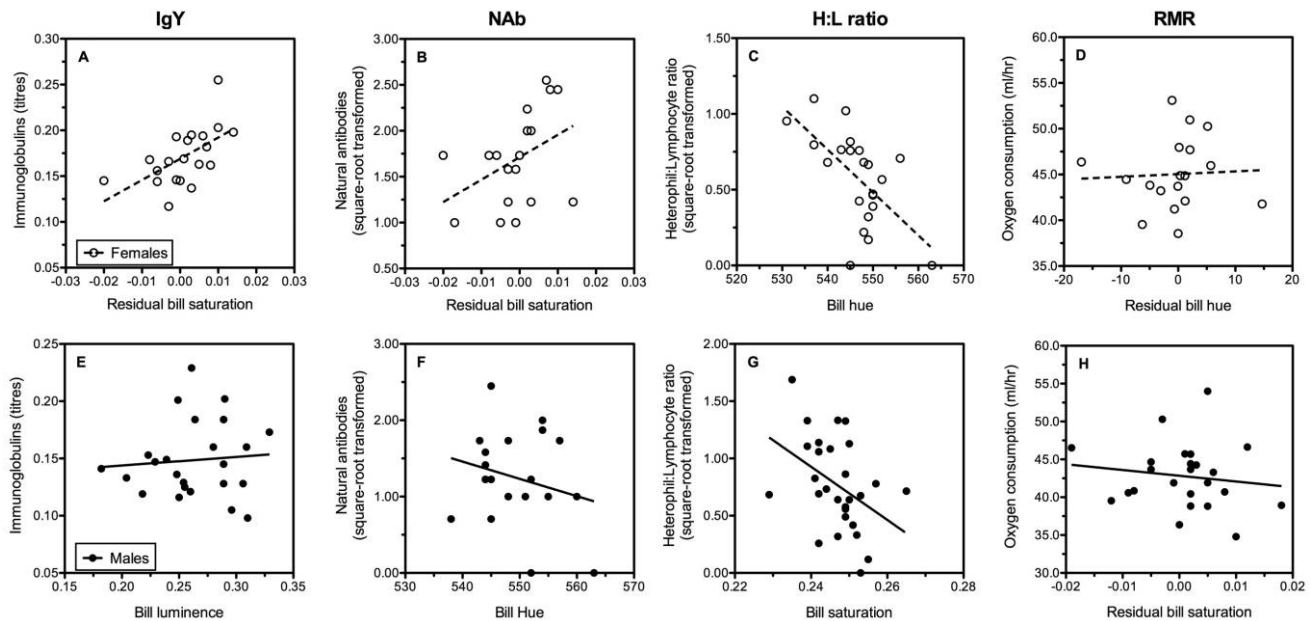


Figure 1. Bill coloration in relation to physiology and immunology of female and male American goldfinches. The independent variable represents the best model of the set that contained a color variable. Note that the best model may have had similar support to a model containing only the intercept (see text for description of best models). Fitted lines are to aid in visual interpretation of patterns. Figures show residuals when the top model included two or more independent variables. Residuals in A and B controlled for the effect of capture date, while D and H controlled for the effect of body mass on the color variable of interest. IgY = immunoglobulin Y; NAb = natural antibodies; H : L ratio = heterophil-to-lymphocyte ratio; RMR = resting metabolic rate.

Table 2: 95% confidence set of best-ranked regression models explaining variation in female bill color

Model	<i>K</i>	AICc	ΔAICc	<i>W<sub>i</sub></i>	Acc <i>W<sub>i</sub></i>	ER
IgY:						
Saturation + date	4	-141.567	.000	.395	.395	...
Saturation	3	-140.639	.928	.248	.643	1.59
Saturation + mass + date	5	-137.964	3.603	.065	.708	6.06
Saturation + mass	4	-137.640	3.927	.055	.763	7.13
Luminance	3	-137.433	4.134	.050	.813	7.90
Intercept only	2	-137.404	4.163	.049	.862	8.02
Luminance + date	4	-136.784	4.783	.036	.898	10.93
Date	3	-135.713	5.854	.021	.920	18.67
Mass	3	-135.247	6.321	.017	.936	23.58
NAb:						
Saturation + date	4	-25.821	.000	.335	.335	...
Date	3	-24.156	1.665	.146	.481	2.30
Mass	3	-23.040	2.781	.083	.564	4.02
Saturation + mass + date	5	-22.708	3.113	.071	.635	4.74
Date + mass	4	-22.316	3.505	.058	.693	5.77
Intercept only	2	-22.188	3.633	.054	.747	6.15
Luminance + date	4	-21.969	3.852	.049	.796	6.86
Hue	3	-21.693	4.128	.043	.839	7.88
Saturation + mass	4	-21.066	4.755	.031	.870	10.78
Hue + date	4	-20.814	5.007	.027	.897	12.23
Hue + mass	4	-20.792	5.029	.027	.924	12.36
Saturation	3	-20.657	5.164	.025	.950	13.22
H : L ratio:						
Hue	3	-58.341	.000	.467	.467	...
Hue + mass	4	-57.188	1.154	.262	.730	1.78
Hue + date	4	-55.504	2.838	.113	.843	4.13
Hue + mass + date	5	-54.539	3.803	.070	.912	6.69
Date	3	-52.156	6.186	.021	.934	22.04
Luminance + date	4	-51.417	6.925	.015	.948	31.89
RMR:						
Mass	3	37.030	.000	.484	.484	...
Date + mass	4	39.271	2.241	.158	.642	3.07
Hue + mass	4	40.179	3.149	.100	.742	4.83
Saturation + mass	4	40.325	3.295	.093	.835	5.19
Luminance + mass	4	40.383	3.352	.091	.926	5.34

Note. IgY = immunoglobulin Y; NAb = natural antibodies; RMR = resting metabolic rate; H : L ratio = heterophil-to-leukocyte ratio; *K* = parameters in the model; AICc = corrected Akaike Information Criterion; ΔAICc = difference in AICc score between focal model and best model; *W<sub>i</sub>* = Akaike weight; Acc *W<sub>i</sub>* = accumulated Akaike weight; ER = evidence ratio.

and the mean IgY absorbance value was used in analysis. The average coefficient of variation (CV) between duplicate samples was 4.96%, and no sample with a CV of more than 15 was used in analysis. IgY absorbance values were obtained for 44 individuals (24 males, 20 females).

#### *H : L Ratio*

Two blood smears were made within 1 h of capture for each individual. Blood smears were stained using Diff-Quik Stain

Set (Dade Behring, Newark, DE). Heterophils and lymphocytes were counted under oil immersion at 1,000× magnification until 100 leukocytes were identified or 150 fields of a homogeneous monolayer of cells had been examined, whichever came first, following Maney et al. (2008). Because smears were of varying quality, all blood smears were initially ranked blindly (by R.J.K.) based on gross appearance (quality of the shape and staining of the smear) and assigned a score of 1–4 (from low to high quality). When two high-quality blood smears were available, we averaged the H : L ratio for the two slides; oth-

Table 3: Full-model-averaged estimates for each variable in the global model predicting female bill color

Response and predictor variables	$W$	$\tilde{\beta}$	$\text{Var}(\tilde{\beta})$
<b>IgY:</b>			
Intercept	...	-.5255	2.1069
Date	.548	.0013	$1.9 \times 10^{-5}$
Mass	.170	.0004	.0003
Luminance	.104	-.0342	.0833
Saturation	.763	1.6710	9.9833
Hue	.040	$5.6 \times 10^{-5}$	$1.3 \times 10^{-6}$
<b>NAb:</b>			
Intercept	...	-8.0297	632.8369
Date	.703	.0351	.0059
Mass	.304	.0689	.1584
Luminance	.091	-.1684	15.6230
Saturation	.462	11.1598	1,350.5058
Hue	.105	-.0015	.0006
<b>H : L ratio:</b>			
Intercept	...	14.6153	577.5950
Date	.243	.0028	.0009
Mass	.356	-.0347	.0469
Luminance	.027	.0636	1.9431
Saturation	.015	-.0979	20.3558
Hue	.912	-.0256	.0015
<b>RMR:</b>			
Intercept	...	-7.6750	14,746.0040
Date	.230	-.0274	.0653
Mass	.998	4.3035	15.1264
Luminance	.117	.3660	655.9392
Saturation	.116	2.2525	21,274.2910
Hue	.124	.0031	.0248

Note.  $W$  = sum of  $W_i$  across all models that include the variable;  $\tilde{\beta}$  = full-model-averaged regression coefficient;  $\text{Var}(\tilde{\beta})$  = full-model-averaged variance of  $\tilde{\beta}$ .

erwise, we analyzed the slide of the highest quality. The H : L ratio was measured for 50 individuals (28 males, 22 females). Sample sizes were larger for the H : L ratio than for NAb and IgY because blood smears were collected in the field, and blood samples were not always sufficiently large to run all remaining assays.

### RMR

Birds were retrieved from the outdoor aviary each night at ca. 1900 hours and transported (<400 m) in a cloth bird bag to a separate laboratory for oxygen consumption trials. Birds were weighed before being placed into metabolic chambers and reweighed at the end of the trial the next morning on removal from the chamber. We report the average mass in subsequent analysis of oxygen consumption.

We measured the overnight oxygen consumption in post-absorptive individuals at thermoneutrality using an open-flow, push-through respirometry system (Sable Systems, Las Vegas,

NV; following Burness et al. 2010). Metabolic chambers containing individual birds were housed within a portable temperature-controlled cabinet (PTC-1, Sable Systems) maintained at 30°C (thermoneutrality; Marsh and Dawson 1982) using a temperature controller (Pelt-5, Sable Systems).

A typical recording sequence consisted of 15 min of measuring oxygen consumption from a chamber containing a bird followed by 5 min of baseline. This sequence continued overnight, alternating among each of the three chambers, from approximately 1945 hours each night to 0530 hours the next morning. Thus, when three birds were simultaneously tested, one 15-min measurement of each bird was obtained every hour for the entire sampling period. To ensure birds were post-absorptive, the first 4 h of O<sub>2</sub> measurements were not considered, and thus ca. 1 h of oxygen consumption measurements were obtained per bird each night. RMR was considered to be the lowest stable 5-min period of oxygen consumption for each bird and was calculated using equation (10.1) from Lighton (2008). RMR was calculated for 42 individuals (24 males, 18 females). The precision of the respirometry system was 97%, estimated by burning a known mass of methanol daily in a metabolic chamber and comparing it with theoretically predicted values.

### Statistical Analyses

All statistical analyses were completed using Statistica, version 7.0 (StatSoft), and PASW Statistics 18.0 (SPSS). We used generalized linear models to directly compare bill and plumage color and physiological parameters between males and females. Analyses of color included Julian capture date as a covariate; analyses of physiological parameters included capture date and mass as covariates.

Following Garamszegi et al. (2009) and Garamszegi (2011), we use an information theoretic approach to infer relationships between color and physiological parameters. Our objective was to determine whether bill and/or plumage color predicted immunological (IgY, NAb) and physiological (H : L ratio, RMR) condition. Because of different function of the signals associated with ornamentation in each sex (see Murphy et al. 2009), males and females were analyzed separately. Likewise, because part of our objective was to test whether dynamic (bill) and static (plumage) ornaments signaled different information, we analyzed those ornaments separately. We assumed a priori that capture date and body mass were likely to influence some or all of our response variables, so we included them as predictor variables in each global model. For each response variable we tested 16 candidate linear regression models that included the following predictor variables either singly or in combination: date, mass, luminance, saturation, and hue or only an intercept. To avoid overfitting models, we included no more than one tristimulus color variable in any model. Thus, the 16 candidate models represent a subset of the possible models that would exist in a full-factorial design (i.e., all combinations of all five variables). We used Akaike Information Criterion-corrected (AICc) scores to assess models because among the models

Table 4: 95% confidence set of best-ranked regression models explaining variation in male bill color

Model	$K$	AICc	$\Delta$ AICc	$W_i$	Acc $W_i$	ER
IgY:						
Intercept only	2	-162.480	.000	.331	.331	...
Date	3	-160.263	2.216	.109	.440	3.03
Luminance	3	-160.015	2.465	.097	.537	3.43
Saturation	3	-159.939	2.541	.093	.630	3.56
Mass	3	-159.910	2.570	.092	.721	3.61
Hue	3	-159.853	2.627	.089	.810	3.72
Luminance + date	4	-157.457	5.023	.027	.837	12.33
Hue + date	4	-157.456	5.024	.027	.864	12.33
Date + mass	4	-157.429	5.051	.026	.890	12.50
Saturation + date	4	-157.415	5.065	.026	.917	12.59
Luminance + mass	4	-157.148	5.331	.023	.940	14.38
NAb:						
Date	3	-16.031	.000	.226	.226	...
Intercept only	2	-15.725	.306	.194	.420	1.17
Mass	3	-13.987	2.044	.081	.501	2.78
Hue	3	-13.940	2.091	.079	.581	2.84
Saturation	3	-13.558	2.473	.066	.647	3.44
Date + mass	4	-13.459	2.572	.062	.709	3.62
Luminance + date	4	-12.960	3.071	.049	.758	4.64
Saturation + date	4	-12.955	3.076	.049	.806	4.66
Hue + date	4	-12.909	3.122	.047	.854	4.76
Luminance	3	-12.894	3.136	.047	.901	4.80
Hue + mass	4	-11.826	4.205	.028	.929	8.19
H : L ratio:						
Saturation	3	-52.437	.000	.395	.395	...
Saturation + mass	4	-49.975	2.462	.115	.511	3.42
Saturation + date	4	-49.700	2.736	.101	.612	3.93
Luminance	3	-49.687	2.749	.100	.712	3.95
Intercept only	2	-49.512	2.925	.092	.803	4.32
Mass	3	-47.368	5.069	.031	.835	12.61
Mass + luminance	4	-47.276	5.160	.030	.865	13.20
Date	3	-47.001	5.436	.026	.891	15.15
Hue	3	-46.992	5.444	.026	.917	15.21
Saturation + mass + date	5	-46.987	5.449	.026	.943	15.25
RMR:						
Mass	3	62.581	.000	.418	.418	...
Saturation + mass	4	64.663	2.082	.148	.566	2.83
Hue + mass	4	65.238	2.658	.111	.676	3.78
Date + mass	4	65.256	2.675	.110	.786	3.81
Luminance + mass	4	65.443	2.863	.100	.886	4.18
Hue + mass + date	5	67.266	4.686	.040	.926	10.41

Note. Abbreviations as in table 2.

tested, our sample sizes ( $n$ ) ranged from 18 to 28 and the number of parameters per model ( $k$ ) ranged from 2 to 5; thus  $n/k < 40$  for all analyses. We used full-model averaging to assess effects of the variables in the global model (Symonds and Mousalli 2011). We evaluated the importance of individual variables as predictors of physiological parameters based on the  $\Delta$ AICc and evidence ratio (ER) scores of models that included the

variable of interest as well as the number of top models that the variable appeared in and the summed Akaike weight ( $W$ ) and the model-averaged slope ( $\hat{\beta}$ ) with its associated variance ( $\text{Var}[\hat{\beta}]$ ) for that variable based on all models in the global set. We paid particular attention to the  $\Delta$ AICc and ER scores of models that included only an intercept. Intercept-only models with low values for these parameters (e.g.,  $< 2$ ) indicate that

Table 5: Full-model-averaged estimates for each variable in the global model predicting male bill color

Response and predictor variables	$W$	$\tilde{\beta}$	$\text{Var}(\tilde{\beta})$
<b>IgY:</b>			
Intercept		.1018	2.3596
Date	.232	.0002	$8.1 \times 10^{-6}$
Mass	.201	.0006	.0008
Luminance	.152	.0106	.1356
Saturation	.147	-.0333	2.5155
Hue	.143	$-4.6 \times 10^{-6}$	$3.3 \times 10^{-6}$
<b>NAb:</b>			
Intercept	...	-.0765	1,024.2883
Date	.465	.0202	.0065
Mass	.243	-.0514	.2553
Luminance	.122	-.1048	28.0158
Saturation	.149	-1.9271	1,177.2817
Hue	.165	-.0029	.0018
<b>H : L ratio:</b>			
Intercept	...	4.1391	261.3802
Date	.202	$4.8 \times 10^{-5}$	.0012
Mass	.228	.0136	.0826
Luminance	.163	.5194	19.3961
Saturation	.638	-14.9671	1,928.8120
Hue	.042	$1.8 \times 10^{-6}$	.0002
<b>RMR:</b>			
Intercept	...	-19.1368	30,498.9734
Date	.211	.0169	.0901
Mass	.980	4.3884	38.5732
Luminance	.124	-.4156	914.3803
Saturation	.181	-13.8960	37,440.3627
Hue	.160	.0143	.0611

Note. Abbreviations as in table 3.

none of the models in the candidate set is particularly good at predicting the response variable.

## Results

### Sex Differences

Plumage of male goldfinches was significantly brighter and more saturated in the yellow region and had greater hue than that of females (table 1). Male bills were significantly brighter and had greater hue than those of females but did not differ in yellow saturation (table 1).

Males and females also differed in measures of immunity (table 1). When controlling for capture date, females had significantly higher NAb titers and IgY absorbance values than males, but males and females did not differ in the H : L ratio. After accounting for differences in body mass, there was no significant sex difference in RMR (table 1).

### Bill Color

In females, bill color was a strong predictor of IgY and the H : L ratio but not other measures of immune function or RMR (fig. 1; tables 2, 3). The positive correlation between bill saturation and IgY was supported by the fact that saturation appeared in the top four models and by the fact that  $W$  for saturation was 0.763, which indicates a probability of 76.3% that it was a component of the true best model (tables 2, 3). Moreover, the ER indicated that the best model of the set, which included saturation and capture date, was 7.9 times better than the next best model that did not include saturation.

Female bill hue was negatively correlated with the H : L ratio (tables 2, 3). The model containing only hue as a predictor variable was the best model in the set, and this model was 22 times better than the next best model that did not contain hue. Moreover,  $W$  for hue was 0.91. We also found some support for a positive relationship between female bill saturation and NAb ( $W = 0.46$ ), although the best model of the set, which included saturation and date, was only about 2.3 times better than a model that included only date, and the  $\Delta\text{AICc}$  score was 1.7 (tables 2, 3).

Mass was the most important predictor of RMR, but models including mass and one of the color variables had weak support ( $W$  for each color variable was approximately 0.12; table 3). However, the model containing only mass garnered approximately five times the evidence of models including mass and any one of the three color variables (table 2).

In males, bill color predicted the H : L ratio but not other measures of immune function or RMR (fig. 1; tables 4, 5). Bill saturation was negatively correlated with the H : L ratio, and saturation appeared in the top three models of the set. The best model of the set contained only saturation, and this model had four times the evidence of the highest-ranked model that did not contain saturation. Additionally,  $W$  for saturation was 0.64. However, we found little or no support for a correlation between bill color and IgY or NAb (tables 4, 5). As with females, mass was the most important predictor of RMR in males, and we found weak support for correlations between color components and RMR when mass also was included as a predictor variable (tables 4, 5).

### Plumage Color

In contrast to bill color, female plumage color predicted NAb but was unrelated to IgY and the H : L ratio (fig. 2; tables 6, 7). Throat hue was positively correlated with NAb: hue appeared in the top two models of the set, and  $W$  for hue was 0.50 (tables 6, 7). The top-ranked model contained hue, mass, and date and was 3.2 times better than the highest-ranking model that did not contain a color variable (date only;  $\Delta\text{AICc} = 2.3$ ).

There was some support for a positive correlation between saturation and NAb as well ( $W = 0.22$ ). We found no support for a relationship between female throat color and IgY or the H : L ratio; models including these variables had ERs that were



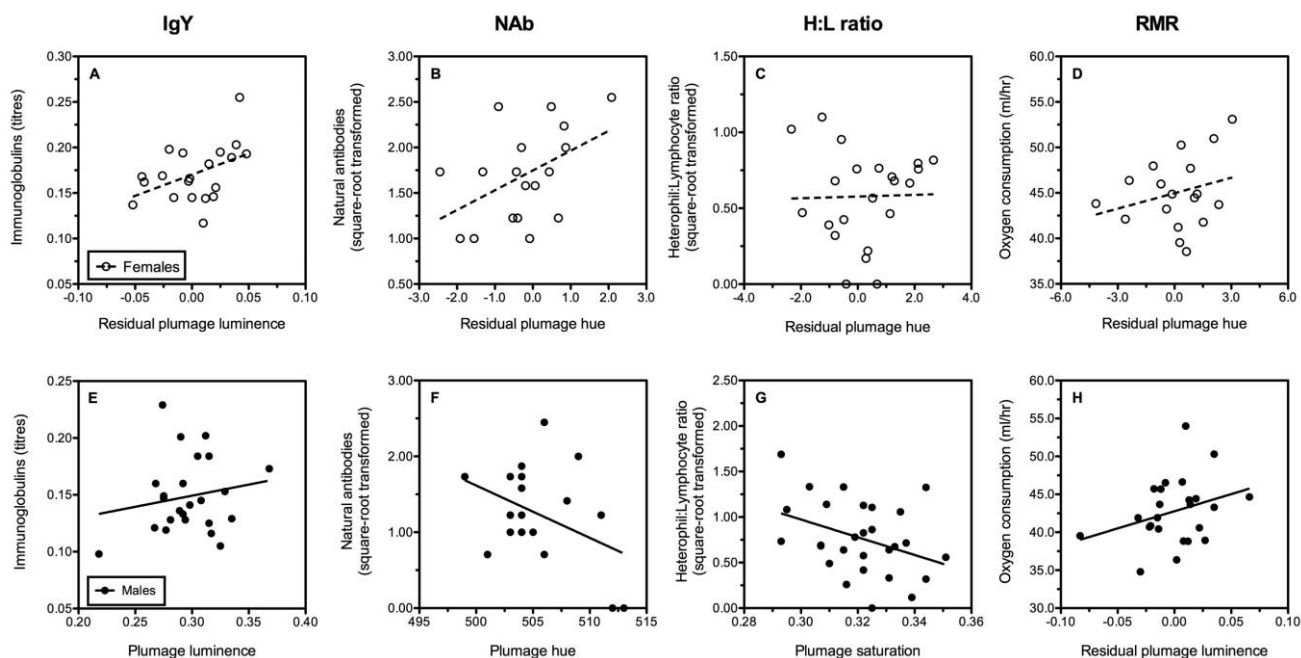


Figure 2. Plumage coloration in relation to physiology and immunology of female and male American goldfinches. The independent variable represents the best model of the set that contained a color variable. Note that the best model may have had similar support to a model containing only the intercept (see text for description of best models). Fitted lines are to aid in visual interpretation of patterns. Figures show residuals when the top model included two or more independent variables. Residuals in *A* and *C* controlled for the effect of date, in *B* controlled for mass and date, and in *D* and *H* controlled for the effect of body mass on the color variable of interest. IgY = immunoglobulin Y; NAb = natural antibodies; H : L ratio = heterophil-to-lymphocyte ratio; RMR = resting metabolic rate.

either equal to or worse than a model that contained only the intercept (table 6).

In the analysis of RMR, mass appeared in each of the models that made up the 95% confidence set, and each color variable appeared in one of the top four models in the set. Of the three color variables, hue had the most support, and the top-ranked model (which contained only mass) had only 1.57 times the evidence of the model that contained both mass and hue. Thus, hue seems to be a predictor of female RMR such that females with high plumage hue had high RMR for their body mass (tables 6, 7; fig. 2).

Male throat color did not predict any of the immunological variables but was a good predictor of RMR (fig. 2; tables 8, 9). Regarding immunological variables, in all cases, even when a color variable had reasonable support relative to other variables in the global model, support for models containing a color variable was similar to that for a model containing only the intercept (i.e.,  $\Delta\text{AICc}$  scores were  $<2.0$ ; tables 8, 9). For RMR, luminance occurred in the top-ranked model (along with mass), and the *W* score was 0.61. Individuals with high luminance had relatively high RMR for their body mass. Even so, the top-ranked model garnered only twice the evidence of a model that included mass but no color variable ( $\text{ER} = 2.02$ ; tables 8, 9).

## Discussion

In general, our findings are consistent with the hypothesis that carotenoid-based bill and plumage colors of American goldfinches function as condition-dependent signals. We showed that ornaments shared by the sexes are related to different aspects of immunity and physiology in males and females. We found that female bill color is strongly positively correlated with adaptive immunity (IgY) and that female plumage color (and to a lesser extent female bill color) is positively correlated with the innate immune system (NAb). However, male ornamentation was not related to either of these components of immunity, thus indicating that sex-specific mechanisms may mediate these signals. We additionally found that males and females with more colorful bills had a lower H : L ratio and that plumage color in both sexes positively reflects RMR, although the specific component of color that correlated with measures of immunity and physiology varied with sex.

Taken together, our results indicate that dynamic bill and static plumage coloration differ in their potential to provide information about short-term versus long-term aspects of physiological condition. The highly variable bill color of both sexes is related to the H : L ratio, which itself is known to fluctuate over the short term (on the order of hours to days) as heterophils increase and lymphocytes decrease in response to stress

Table 6: 95% confidence set of best-ranked regression models explaining variation in female throat color

Model	<i>K</i>	AICc	$\Delta$ AICc	$W_i$	Acc $W_i$	ER
IgY:						
Luminance + date	4	-137.448	.000	.175	.175	...
Intercept only	2	-137.404	.045	.171	.347	1.02
Luminance	3	-136.864	.584	.131	.477	1.34
Luminance + mass + date	5	-136.803	.645	.127	.604	1.38
Luminance + mass	4	-135.883	1.565	.080	.684	2.19
Date	3	-135.713	1.735	.074	.758	2.38
Mass	3	-135.247	2.202	.058	.816	3.01
Hue	3	-135.148	2.301	.055	.872	3.16
Saturation	3	-134.611	2.837	.042	.914	4.13
Date + mass	4	-133.063	4.386	.020	.934	8.96
NAb:						
Hue + mass + date	5	-26.477	.000	.308	.308	...
Hue + date	4	-25.259	1.218	.168	.476	1.84
Saturation + date	4	-24.464	2.013	.113	.589	2.74
Date	3	-24.156	2.321	.097	.685	3.19
Saturation + mass + date	5	-23.739	2.738	.078	.764	3.93
Mass	3	-23.040	3.437	.055	.819	5.58
Date + mass	4	-22.316	4.161	.038	.857	8.01
Intercept only	2	-22.188	4.289	.036	.894	8.54
Hue + mass	4	-21.046	5.431	.020	.914	15.11
Luminance + date	4	-20.893	5.584	.019	.933	16.32
H : L ratio:						
Date	3	-52.156	.000	.260	.260	...
Date + mass	4	-51.117	1.039	.155	.414	1.68
Intercept only	2	-50.428	1.728	.110	.524	2.37
Hue + date	4	-49.437	2.718	.067	.591	3.89
Luminance + date	4	-49.151	3.005	.058	.649	4.49
Saturation + date	4	-49.147	3.009	.058	.706	4.50
Saturation	3	-48.920	3.236	.052	.758	5.04
Hue	3	-48.521	3.635	.042	.800	6.16
Luminance	3	-48.266	3.889	.037	.837	6.99
Mass	3	-48.027	4.129	.033	.870	7.88
Luminance + mass + date	5	-47.870	4.286	.030	.901	8.52
Hue + mass + date	5	-47.850	4.306	.030	.931	8.61
RMR:						
Mass	3	37.030	.000	.338	.338	...
Hue + mass	4	37.934	.903	.215	.554	1.57
Luminance + mass	4	38.909	1.879	.132	.686	2.56
Saturation + mass	4	39.052	2.021	.123	.809	2.75
Date + mass	4	39.271	2.241	.110	.919	3.07

Note. Abbreviations as in table 2.

(Davis et al. 2008). In contrast, plumage color in both sexes, which is stable over periods of time on the order of months, reflects RMR, which is likely consistent within individuals over time. Although RMR is known to be influenced by embryonic (Nilsson et al. 2011) and neonatal (Verhulst et al. 2006; Criscuolo et al. 2008) environments, significant repeatability in adult RMR has been reported over multiple years in the lab and field (Rønning et al. 2005; Broggi et al. 2009, reviewed by

Biro and Stamps 2010). If RMR is similarly relatively stable within adult goldfinches during the months before the nesting period, then our results may reflect a relationship between plumage color and RMR that persists several months after the prealternate moult. Furthermore, female plumage also reflects NAb levels. Constitutive innate NAb levels appear to vary little with environmental influences and are germ-line encoded and therefore likely reflect long-term (genetic) investment in innate

Table 7: Full-model-averaged estimates for each variable in the global model predicting female throat color

Response and predictor variables	$W$	$\tilde{\beta}$	$\text{Var}(\tilde{\beta})$
<b>IgY:</b>			
Intercept	...	-.0214	9.7076
Date	.435	.0011	$1.3 \times 10^{-5}$
Mass	.318	.0036	.0006
Luminance	.513	.2341	.5536
Saturation	.073	.0024	.4650
Hue	.091	-.0002	$3.0 \times 10^{-5}$
<b>NAb:</b>			
Intercept	...	-64.2750	22,361.6500
Date	.828	.0524	.0082
Mass	.536	.1584	.2484
Luminance	.052	-.0394	18.3249
Saturation	.216	3.6149	376.4613
Hue	.505	.1046	.0778
<b>H : L ratio:</b>			
Intercept	...	-2.4982	2,084.7220
Date	.686	.0181	.0025
Mass	.317	-.0332	.0588
Luminance	.137	-.1256	16.7791
Saturation	.154	-.3001	85.2596
Hue	.153	$2.8 \times 10^{-5}$	.0070
<b>RMR:</b>			
Intercept	...	-70.7990	143,776.1000
Date	.191	-.0179	.0509
Mass	.999	4.3936	13.6359
Luminance	.158	4.5956	2,116.2750
Saturation	.143	7.1490	6,727.6730
Hue	.249	.1177	.5319

Note. Abbreviations as in table 3.

immunity (Tieleman et al. 2005). Thus, our results suggest that in goldfinches, the speed at which bill and plumage signals change over time in general corresponds with the speed at which change occurs in the component of physiology that is reflected by each ornament. These results suggest that dynamic and static signals provide information about different temporal aspects of condition (the “multiple messages” hypothesis; Møller and Pomiankowski 1993).

Our finding that plumage color is positively correlated with RMR in both sexes is particularly interesting because we are unaware of any study that has correlated carotenoid coloration with metabolic rate. High RMR is thought to increase the capacity to perform well in challenging activities, and individuals with high RMR tend to have higher growth rates, have higher activity levels, and be more dominant (reviewed in Biro and Stamps 2010). This is likely to translate into a higher capacity to perform behavioral displays; provision young; and compete for food, mates, and nest sites, and receivers may benefit from identifying potential competitors or mates that have these qualities. On the other hand, it has been argued that individuals

with high RMRs have higher activity rates and resulting oxidative damage (Wiersma et al. 2004). Accordingly, individuals with high RMRs would require more carotenoids to prevent oxidative damage. Given this likely trade-off, we expect individuals in good condition to be better able to sustain an elevated RMR (so as to increase the “capacity to engage in costly behaviour”; Biro and Stamps 2010, p. 657) while also producing colorful ornaments. Thus, male and female goldfinches with colorful plumage appear to be able to sustain the benefits of high RMR while also sustaining highly functional immune systems and colorful ornamentation.

Interpretation of the relationship between ornamentation and adaptive immunity as indicated by IgY levels is difficult because high antibody levels may reflect high immune capacity (via constituent antibodies that are ready to engage with pathogens) or an induced response to a specific infection agent (see Morales et al. 2004 for discussion of this dilemma). Thus, in the first case we would expect a positive correlation between an ornamental trait and IgY to represent a signal of high condition, as individuals with high levels of IgY should be highly immunocompetent. In contrast, in the second case a positive correlation would suggest that highly ornamented individuals were more susceptible to or more likely exposed to infection because they are likely to be fighting a specific infection at the time of blood sampling. To help discern between these possibilities, we can look at the distribution of IgY across individuals. Individuals within a population mounting an adaptive response to a current infection should have substantially higher antibody levels than individuals that are not engaged in fighting a particular infection. If IgY level indicates current infection, we would expect the distribution of antibody levels to be abruptly skewed because only some individuals in a population would be expected to be fighting an infection at a given time, and those individuals should have much higher levels of IgY than individuals that are not fighting an infection (see Butler et al. 2010; Ferreira et al. 2010). However, the observed distribution of IgY in goldfinches is monotonic (fig. 1A, 1E), suggesting that our measure of IgY is a measure of constituent antibodies and reflects individual differences in immune preparedness.

Our observation that highly ornamented females have stronger immune systems (i.e., higher levels of both IgY and NAb) suggests that these females were in particularly good condition at the time of sampling (as bill color reflects condition-dependent IgY) and during the lengthy moult period that occurred several weeks before sampling (as plumage color reflects genotype-dependent NAb). This relationship is consistent with the hypothesis that there is a direct trade-off between use of carotenoids in ornamentation and immunity or oxidative stress management and that only high-quality individuals can successfully allocate carotenoids to both ornamentation and somatic maintenance. Alternatively, these results are consistent with the hypothesis that the degree of ornamentation and immune function each are dependent on the quality of subcellular and biochemical processes such that individuals in good condition (i.e., in which these processes function well) are simultaneously able to generate colorful carotenoid-based orna-

Table 8: 95% confidence set of best-ranked regression models explaining variation in male throat color

Model	<i>K</i>	AICc	$\Delta$ AICc	$W_i$	Acc $W_i$	ER
IgY:						
Intercept only	2	-162.480	.000	.303	.303	...
Luminance	3	-160.598	1.882	.118	.422	2.56
Date	3	-160.263	2.216	.100	.522	3.03
Saturation	3	-160.146	2.334	.094	.616	3.21
Hue	3	-159.944	2.536	.085	.701	3.55
Mass	3	-159.910	2.570	.084	.785	3.61
Luminance + date	4	-158.814	3.666	.048	.834	6.25
Luminance + mass	4	-157.841	4.639	.030	.864	10.17
Saturation + date	4	-157.439	5.041	.024	.888	12.43
Date + mass	4	-157.429	5.051	.024	.912	12.50
Hue + date	4	-157.360	5.120	.023	.936	12.93
NAb:						
Hue	3	-16.201	.000	.172	.172	...
Date	3	-16.031	.170	.158	.329	1.09
Saturation	3	-15.820	.381	.142	.471	1.21
Intercept only	2	-15.725	.476	.135	.606	1.27
Mass	3	-13.987	2.214	.057	.663	3.02
Hue + date	4	-13.862	2.339	.053	.716	3.22
Hue + mass	4	-13.620	2.581	.047	.763	3.63
Saturation + mass	4	-13.522	2.679	.045	.808	3.82
Date + mass	4	-13.459	2.742	.044	.852	3.94
Saturation + date	4	-13.400	2.801	.042	.894	4.06
Luminance	3	-12.922	3.278	.033	.927	5.15
H : L ratio:						
Saturation	3	-51.351	.000	.292	.292	...
Saturation + date	4	-49.556	1.796	.119	.412	2.45
Intercept only	2	-49.512	1.839	.117	.528	2.51
Hue	3	-49.374	1.977	.109	.637	2.69
Saturation + mass	4	-49.181	2.171	.099	.736	2.96
Luminance	3	-47.502	3.850	.043	.778	6.85
Mass	3	-47.368	3.983	.040	.818	7.33
Hue + mass	4	-47.286	4.065	.038	.857	7.63
Date	3	-47.001	4.350	.033	.890	8.80
Saturation + mass + date	5	-46.963	4.388	.033	.922	8.97
RMR:						
Luminance + mass	4	61.176	.000	.402	.402	...
Mass	3	62.581	1.405	.199	.601	2.02
Luminance + mass + date	5	62.613	1.437	.196	.796	2.05
Saturation + mass	4	64.951	3.775	.061	.857	6.60
Date + mass	4	65.256	4.080	.052	.909	7.69
Hue + mass	4	65.262	4.086	.052	.962	7.71
Saturation + mass + date	5	68.158	6.982	.012	.974	32.82

Note. Abbreviations as in table 2.

ments, sustain highly functional immune systems, and deal with oxidative stress (Hill 2011). Whether or not direct trade-offs are involved, the patterns we observed are consistent with the hypothesis that carotenoid-based ornaments signal immunological capacity in female goldfinches. The reasons that such ornaments do not act as a similar signal of immunological

capacity in male goldfinches is not clear, especially given the links between carotenoid-based signals and aspects of immunity in other species (e.g., Aguilera and Amat 2007; Baeta et al. 2008; Dunn et al. 2010).

There is good reason to expect goldfinches to make use of reliable signals of quality in both sexual and competitive con-

Table 9: Full-model-averaged estimates for each variable in the global model predicting male throat color

Response and predictor variables	W	$\tilde{\beta}$	Var( $\tilde{\beta}$ )
<b>IgY:</b>			
Intercept	...	.1247	5.1162
Date	.241	.0002	$1.0 \times 10^{-5}$
Mass	.202	.0007	.0008
Luminance	.208	.0465	.2955
Saturation	.147	-.0318	.7763
Hue	.134	-.0001	$1.3 \times 10^{-5}$
<b>NAb:</b>			
Intercept	...	8.9741	3,014.1640
Date	.356	.0138	.0056
Mass	.233	-.0466	.2396
Luminance	.086	-.1283	46.7869
Saturation	.238	-2.9058	368.9960
Hue	.283	-.0177	.0096
<b>H : L ratio:</b>			
Intercept	...	6.4036	987.5518
Date	.250	-.0025	.0018
Mass	.244	.0187	.0911
Luminance	.069	-.1235	14.2329
Saturation	.543	-5.7172	394.3418
Hue	.188	-.0069	.0032
<b>RMR:</b>			
Intercept	...	-21.5376	28,906.6800
Date	.276	.0326	.0981
Mass	.987	4.2292	34.9743
Luminance	.607	30.3481	8,541.9210
Saturation	.074	-2.2339	3,941.7780
Hue	.064	-.0056	.0749

Note. Abbreviations as in table 3.

texts. Both males and females forage in flocks that vary in composition throughout the year (McGraw and Middleton 2009) and are likely to routinely engage in competitive interactions with both familiar and unfamiliar individuals. Furthermore, both sexes contribute substantially to parental care (McGraw and Middleton 2009), and as a result, both sexes may benefit from assessing quality of potential mates (Johnstone et al. 1996; Amundsen 2000). In our earlier work, we found that female (but not male) bill color functions as a signal to other females during competition for nonsexual resources such as food (Murphy et al. 2009) but that bill color does not appear to be used in mate choice by either sex (Murphy et al. 2009; T. G. Murphy and K. A. Tarvin, unpublished data). Thus, the finding of this study that female bill color is positively correlated with immune function suggests that immunological capacity plays an important role in determining a female's ability to invest in competitive interactions or that perhaps bill color and immunity both represent general measures of condition. Our findings also indicate that both male plumage and female plumage have the potential to function as a signal of quality and

that both sexes may be assessed during mate choice (see MacDougall and Montgomerie 2003). Several studies have shown that male goldfinch plumage color signals general aspects of nutritional and health status (e.g., McGraw and Hill 2000, 2001; Rosen and Tarvin 2006; Hill et al. 2009) and that colorful male plumage is preferred by females (Johnson et al. 1993). Future work is necessary to assess the signaling role of female plumage.

In this study, we identify specific components of immunological and physiological condition that are signaled by both male and female ornaments and show that dynamic and static ornaments signal components of condition that differ in the timescale over which they vary. Our results suggest that ornaments can have a sex-specific functional role in social and sexual interactions even when those traits are expressed similarly in the sexes. Our work thus raises questions about how the social and sexual contexts of signals influence their information content.

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