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Losses of female song with changes from tropical to temperate breeding in the New World blackbirds

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Birds in which both sexes produce complex songs are thought to be more common in the tropics than in temperate areas, where typically only males sing. Yet the role of phylogeny in this apparent relationship between female song and latitude has never been examined. Here, we reconstruct evolutionary changes in female song and breeding latitude in the New World blackbirds (Icteridae), a family with both temperate and tropical representatives. We provide strong evidence that members of this group have moved repeatedly from tropical to temperate breeding ranges and, furthermore, that these range shifts were associated with losses of female song more often than expected by chance. This historical perspective suggests that male-biased song production in many temperate species is the result not of sexual selection for complex song in males but of selection against such songs in females. Our results provide new insights into the differences we see today between tropical and temperate songbirds, and suggest that the role of sexual selection in the evolution of bird song might not be as simple as we think.

Keywords: ancestral state reconstruction; comparative methods; duetting; female bird song; phylogeny; sexual selection

1. INTRODUCTION

Most research on the evolution of bird song has focused on north temperate passerines in which songs are produced almost exclusively by males (Marler & Slabbekoorn 2004; Catchpole & Slater 2008). Female song is rare in temperate species (Riebel 2003), and where it does occur it is produced less frequently than male song and often with less complexity (e.g. Beletsky 1982, 1983; Arcese et al. 1988). In the tropics, however, females of many species sing (Kroodsma et al. 1996; Morton 1996; Langmore 1998; Slater & Mann 2004), either coordinating their vocalizations with males to perform complicated duets (Farabaugh 1982; Hall 2004), or producing solo songs similar to those of males (Whittingham et al. 1992, 1997; Price et al. 2008). Female bird song may even be the norm rather than the exception in tropical environments (Morton 1996), yet we know surprisingly little about the origins of this latitudinal difference (Slater & Mann 2004). Furthermore, the apparent correlation between tropical breeding and female song has never been tested using methods that correct for phylogeny. Without controlling for phylogeny, one or a few species-rich tropical clades with female song could strongly bias our interpretations and create an apparent trend where none actually exists (Felsenstein 1985; Harvey & Pagel 1991).

Hypotheses to explain the greater prevalence of female song in the tropics have generally focused on factors selecting for tropical females to sing as males, such as mate attraction, shared territorial defence or other forms of behavioural coordination by long-term pairs (Langmore 1998, 2000; Hall 2004; Slater & Mann 2004). This focus

may lead some researchers to assume that the ancestors of these birds resembled temperate species in lacking female song, but it is equally plausible that both sexes sang in ancestors and that female singing has been repeatedly lost during the evolution of temperate taxa (Riebel et al. 2005; Garamszegi et al. 2007). Indeed, such losses of female song seem especially probable in temperate species that have recent tropical ancestors (Kondo & Omland 2007; Barker et al. 2008). Rather than asking why females sing in the tropics (e.g. Slater & Mann 2004), it may be more appropriate in many cases to ask why females do not sing in temperate-breeding species. Yet, the idea that selection could favour losses of female song during vocal evolution has received comparatively little attention.

Here we reconstruct the evolution of female song and compare it to breeding latitude in the New World blackbirds (Icteridae), a songbird family with both temperate and tropical representatives (Jaramillo & Burke 1999). The behavioural and ecological traits of many members of this clade have been intensively studied (Bent 1958; Orians 1985; Robinson 1986; Skutch 1996; Jaramillo & Burke 1999; Searcy et al. 1999; Fraga 2008), and some are models for investigating the evolution of bird song by sexual selection (e.g. Agelaius phoeniceus: Searcy & Yasukawa 1995). Moreover, female song rates vary substantially across the family, from species in which both sexes regularly produce similar songs (e.g. Agelaius assimilis: Whittingham et al. 1992, 1997; Icterus pustulatus: Price et al. 2008) or interpose their notes to produce complicated duets (e.g. Dives dives: Orians 1983), to taxa in which female song is relatively rare or absent (e.g. Icterus galbula: Beletsky 1982; Sturnella neglecta: Lanyon 1994). Substantial differences in female song have even been

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reported between closely related sister taxa (A. phoeniceus and A. assimilis: Whittingham et al. 1992; Barker et al. 2008). The New World blackbirds therefore provide an ideal model clade for addressing questions about the evolution of female bird song, including changes that have occurred over relatively short time scales.

Our phylogenetic analysis focused on 65 tropical and temperate blackbird species with well documented vocal behaviours and well established molecular relationships. Relationships among these taxa were based on a larger phylogeny of the entire family Icteridae (Lanyon & Barker 2007), which uses DNA sequence data from multiple mitochondrial and nuclear loci to resolve relationships among the family's four previously defined subclades (Lanyon & Omland 1999): the oropendolas and caciques, the orioles, the grackles and allies, and the meadowlarks and allies. Our main objectives were to examine: (i) whether current differences among taxa in female song are the result of historical gains or losses, (ii) whether or not these changes were associated with changes in breeding latitude, and (iii) whether similar evolutionary patterns have occurred in each of the four major subclades. We hope that the historical patterns revealed in this study will provide a useful starting point for future investigations into the ecological and social pressures selecting for differences in female song.

2. MATERIAL AND METHODS

(a) Scoring female song and breeding latitude

We gathered information on female singing behaviour from species accounts in identification guides and from published studies (listed in appendix A). We defined 'songs' as vocalizations used in defending resources or advertising for mates, following the definitions described by authors of species account descriptions (Howell & Webb 1995; Jaramillo & Burke 1999). Songs of most New World blackbird species are regularly given in association with a visual display, such as the 'bow display' of oropendolas and the 'song spread' of many other icterids (Jaramillo & Burke 1999), so it was presumably straightforward for previous authors to identify male and female vocalizations as songs rather than other types of sounds in these taxa. We did not include species in which song was not clearly described or in which singing differences between the sexes are not known (all excluded taxa were tropical or south temperate and were distributed across 15 genera). Altogether, we scored singing behaviour in 65 of the approximately 101 named species in the family Icteridae (Clements 2007).

Female song can differ from male song in a variety of ways, including rates of production and general acoustic features. In our analysis, we focused solely on the reported rates of female song relative to male song and ignored comparisons of song structure, since most descriptions did not include sound spectrograms and we could not quantify acoustic differences between songs based on verbal descriptions alone. It was also difficult to reliably differentiate between females that sing relatively infrequently and those that never sing, since infrequent female song in some species could have gone unnoticed by researchers (Garamszegi *et al.* 2007). Therefore, we combined these cases (i.e. species reported to have infrequent female song and those in which females are not known to sing) into one discrete character state in our analysis and scored each species as having either: (i) similar song rates

in either sex or (ii) females that sing less often than males. Scoring female song rates as a binary character ignored much of the variation in female singing behaviour across species but this approach also allowed us to reconstruct changes in a way that was biased against identifying ancestors as having female song, since taxa were given this score only if females sing as often as males.

Species scored as having similar male–female song rates included taxa in which both sexes regularly produce solo songs or combine their songs into antiphonal duets. In at least one species (*I. pustulatus*), females sing more often than males during nest-building and following the breeding season (Price *et al.* 2008); we scored this taxon as having similar song rates in the sexes based on reports that males may sing more often during other parts of the year (Jaramillo & Burke 1999). When only male song was described in detail, we assumed that female song was either rare or absent and scored these species as 'females sing less often than males'. An independent observer unfamiliar with the taxa and our hypotheses also scored female song rates based on our compiled literature descriptions to verify that our scoring was accurate.

We determined breeding ranges using range maps in Jaramillo & Burke (1999) and scored each species as having either a (i) tropical, (ii) north temperate or (iii) south temperate breeding ranges. A species was scored as tropical if approximately 80 per cent or more of its breeding range fell between 23.5° latitude north and south and was scored as north or south temperate if at least 80 per cent of its breeding range fell north or south of these latitudes, respectively. If two zones each contained more than 20 per cent of a breeding range's area (north temperate and tropical: five species; south temperate and tropical: six species; see appendix A), we scored the species based on the location in which most previous studies or observations of singing behaviour had been conducted.

(b) Ancestral state reconstructions and comparative analyses

We determined relationships among our 65 taxa using a molecular phylogeny for the entire family Icteridae (Lanyon & Barker 2007). This phylogeny is based on DNA sequence data from two mitochondrial gene regions (cytochrome *b* and ND2) and four nuclear regions (RAG1, beta fibrinogen intron 5, aconitase 1 intron 10 and myoglobin intron 2). Mitochondrial data were obtained for all taxa, whereas nuclear data were obtained from selected divergent taxa (i.e. at least one sample from every genus and multiple samples from within genera with mitochondrial DNA divergences exceeding 7%).

We reconstructed ancestral character states for female song and breeding latitude on the rooted phylogeny using unordered parsimony in MacClade v. 4.06 (Maddison & Maddison 2003) and using maximum-likelihood methods in Mesquite v. 2.5 (Maddison & Maddison 2008). Our maximum-likelihood analyses used the Markov k-state one-parameter model, which assumes equal rates of change between character states. Maximum likelihood has advantages over parsimony in that it uses information about amounts of genetic divergence between taxa (i.e. branch lengths) and indicates degree of uncertainty in possible ancestral states. In all of these reconstructions, we refer to decreases in the occurrence of female song relating to male song as 'losses' and increases in female song as 'gains'.

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We tested for an association between changes in female song and breeding latitude using the concentrated changes test in MacClade (Maddison 1990). This parsimony-based approach requires each character under comparison to have two states, so we converted breeding latitude from a multistate to a binary character by combining north temperate and south temperate ranges into one character state. We felt justified in combining these states because only 1 out of the 18 changes in our multi-state parsimony reconstruction of breeding range was between north and south temperate ranges (see below). We used breeding latitude as our independent character in this analysis to test the hypothesis that losses of female song have been concentrated in temperate lineages. When a character had more than one most parsimonious reconstruction, we performed multiple tests using either DELTRAN (that favours convergent changes in character states over reversals) or ACCTRAN (that favours reversals over convergent changes) assumptions for each character. Each test included 10 000 simulations.

We further examined the relationship between changes in female song and breeding latitude using Pagel's (1994) discrete likelihood correlation method, as implemented in MESQUITE (Maddison & Maddison 2008). This likelihood-based method tests for an association between discrete character state changes in two binary characters by comparing a dependent model of correlated trait evolution to an independent model of uncorrelated character state changes. We calculated the likelihood of each model and relative transition rate parameters between each possible pair of character states using 10 000 simulation replicates.

3. RESULTS

(a) Reconstructing ancestral states

Parsimony reconstructions of female song evolution (figure 1a) suggest that female song rates have decreased repeatedly in comparison to male song in the New World blackbirds. Sixteen most parsimonious reconstructions were possible on the tree, and all showed multiple independent losses of female song (14 losses and no gains in DELTRAN reconstruction; 11 losses and three gains in ACCTRAN reconstruction). All reconstructions indicate that males and females had similar song rates in the ancestors of the oropendolas and caciques, the orioles and the grackles and allies, and that females sang less than males in the ancestor of the meadowlarks and allies.

Parsimony reconstructions of breeding latitude (figure 1b) showed that taxa have moved from tropical to temperate breeding areas repeatedly, much more often than the reverse. All 16 possible reconstructions agree that ancestors of the oropendolas and caciques, orioles and grackles and allies had tropical breeding ranges, and that the ancestor of the meadowlarks and allies was north temperate. More movements have occurred to and from north temperate areas (11) than south temperate areas (6), using either DELTRAN or ACCTRAN assumptions, which makes sense given the greater land area in the north. We reconstructed only one direct exchange between north and south temperate ranges, and this may be an artefact of scoring a species that is widely distributed in the tropics (Molothrus bonariensis) as a south temperate breeder because most observations of its vocal behaviour have been in the south temperate zone (Jaramillo & Burke 1999; Lowther & Post 1999). Neither breeding range nor female song frequency was resolved for the ancestor of the family.

Maximum-likelihood reconstructions of female song (figure 2a) provided mixed support for the patterns suggested by parsimony reconstructions. Likelihood values agreed with parsimony in strongly indicating that males and females had similar song rates in the ancestor of the orioles (likelihood=0.80) and that females sang less than males in the ancestor of the meadowlarks and allies (0.70). However, unlike in parsimony reconstructions, likelihood values for the ancestors of the oropendolas and caciques, and the grackles and allies provided little support for the idea that males and females in these taxa had similar song rates (0.36 and 0.34, respectively). The only changes in female song that were clearly supported by both reconstruction methods were in the oriole clade, in which losses have occurred at least three times independently.

By contrast, maximum-likelihood reconstructions of breeding latitude (figure 2b) largely agreed with parsimony reconstructions by indicating tropical breeding ranges for ancestors of the oropendolas and caciques (likelihood=0.91), orioles (0.93) and grackles and allies (0.91). Tropical breeding was also suggested for the ancestor of the meadowlarks and allies (likelihood=0.60) and the ancestor of the Icteridae (0.78). Thus, both reconstruction methods showed repeated historical movements by icterids from tropical to temperate breeding ranges.

(b) Correlated evolution between female song and breeding latitude

Concentrated change tests using all possible combinations of DELTRAN and ACCTRAN assumptions supported a strong and significant association between temperate breeding and decreases in the occurrence of female song, with all tests producing p-values of less than 0.01 (DELTRAN song versus DELTRAN latitude: p = 0.0006; ACCTRAN song versus DELTRAN latitude: p=0.0091; DELTRAN song versus ACCTRAN latitude: p=0.0005; ACCTRAN song versus ACCTRAN latitude: p = 0.0001). Of the 11 or more independent losses of female song shown in parsimony reconstructions, at least seven have occurred on branches of the tree in which temperate breeding is the reconstructed state (figure 1). Furthermore, these losses have occurred in both north temperate and south temperate taxa. Only two temperate-breeding icterid species are reported to have frequent female song: Icterus bullockii in the north (Miller 1931; Rising & Williams 1999) and Cacicus chrysopterus in the south (Jaramillo & Burke 1999).

A relationship between changes in female song and breeding latitude was also supported using Pagel's (1994) test for correlated evolution. Comparisons of maximum-likelihood models indicated that the dependent model of correlated trait evolution (log likelihood=-67.9) was significantly more likely than the independent model (log likelihood=-76.2), with a likelihood ratio of 8.3 (p < 0.05). Estimated transition rates (q) between each combination of character states in the dependent model showed that losses of female song were far more likely to occur in temperate breeders (q31=319.1) than tropical breeders (q42=0.00), while gains in the occurrence of female song were relatively unlikely in all lineages (q13=0.00, q24=0.13).

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Figure 1. Ancestral state reconstructions of (a) female song rates and (b) breeding range latitudes calculated using unordered parsimony on a molecular phylogeny of the New World blackbird family. Female song rates were scored as a binary character relative to male song rates and breeding ranges were assessed relative to 23.5° latitude north and south. Four major subclades within the family are indicated on the right. These reconstructions indicate that female song rates have decreased repeatedly with movements from tropical to temperate breeding ranges. (a) Black bars, similar song rates in both sexes; white bars, females sing less than males; striped bars, equivocal. (b) Black bars, tropical breeding; white bars, north temperate breeding; grey bars, south temperate breeding; striped bars, equivocal.

Notable exceptions to the association between female song and latitude can be seen in the oropendolas and caciques, in which female song has changed multiple times among tropical taxa, and in the cowbird clade (genus *Molothrus*), in which parsimony suggests that female song was lost in a tropical ancestor (figure 1). Likewise, where reversals from temperate to tropical breeding ranges have occurred in this family, as appears to have happened in

Chrysomus icterocephalus and twice in the genus Quiscalus, these movements have not been accompanied by apparent changes in female song.

4. DISCUSSION

The view that female song is more common in tropical than in temperate songbirds has been widely held among researchers (Kroodsma *et al.* 1996; Morton 1996;

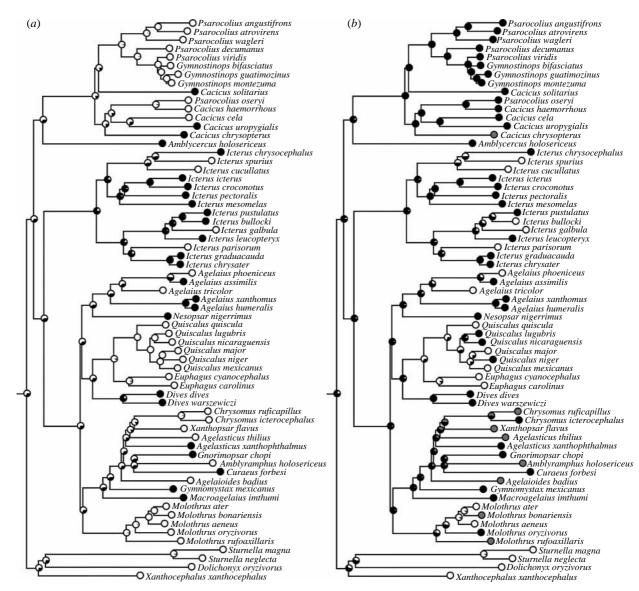


Figure 2. Maximum-likelihood reconstructions of (a) female song rates and (b) breeding range latitudes, assuming equal rates of change between characters states (Markov k-state one parameter model). Black and white fill in circles indicate the proportional likelihoods of ancestral states. (a) Black dots, similar song rates in both sexes; white dots, females sing less than males. (b) Black dots, tropical breeding; white dots, north temperate breeding; grey dots, south temperate breeding.

Langmore 1998; Slater & Mann 2004). To our knowledge, however, our study is the first to test this association using a phylogenetic approach. By controlling for phylogeny, we show that differences in female song between tropical and temperate New World blackbird species are not explained by just a few species-rich clades (Felsenstein 1985) but rather are the result of repeated, correlated historical changes in female song and breeding latitude. Frequent singing by females is significantly associated with tropical breeding in this family; a relationship supported using multiple phylogenetic comparative methods. Females of temperate breeding species almost invariably sing less often than males, if at all, and evolutionary reconstructions show that most of these temperate species had tropical ancestors. Our results therefore suggest that female song has been lost repeatedly in this group in association with repeated evolutionary movements from tropical to temperate breeding areas, and further that these changes have occurred in both the northern and southern hemispheres.

The relationship between losses of female song and increases in latitude is particularly striking in the orioles,

in which both parsimony and maximum-likelihood reconstructions clearly indicate that ancestral taxa were tropical with similar song rates in males and females (figures 1 and 2). Losses of female song have occurred at least three separate times in this clade, and each case was associated with a change from tropical to temperate breeding. Previous reconstructions of migratory behaviours in orioles (Kondo & Omland 2007) show that migration evolved rapidly and relatively recently in these same species (Icterus spurius, Icterus cucullatus, I. galbula and Icterus parisorum). Kondo & Omland (2007) sampled more oriole taxa in their study than we include here (45 taxa from 25 species, rather than the 14 species here), and if losses of female song accompanied the evolution of migration to temperate areas, as these studies together suggest, such losses have occurred four separate times in the orioles and over relatively short time periods. These same lineages have lost elaborate female coloration (Hofmann et al. 2008) that may have also happened in parallel with the evolution of temperate breeding and migration.

Further evidence for rapid evolutionary changes in female song is provided by comparing A. phoeniceus, the

red-winged blackbird of North America and its closest relative A. assimilis, the red-shouldered blackbird of Cuba. These sister taxa are very close genetically and in fact have been considered conspecific in the past (Barker et al. 2008), yet they differ in a variety of morphological and behavioural characteristics (Whittingham et al. 1992). Females of A. assimilis are nearly indistinguishable from conspecific males in song structure and song rate and are also similar in plumage and body size (Whittingham et al. 1992, 1996, 1997; Jaramillo & Burke 1999), whereas females of A. phoeniceus differ considerably from conspecific males in these traits (Beletsky 1983; Searcy & Yasukawa 1995; Yasukawa & Searcy 1995). Whether these male-like female characteristics were lost in A. phoeniceus or gained in A. assimilis was not resolved in our reconstructions of song (figures 1a and 2a) or in similar reconstructions of other traits in these birds (Barker et al. 2008). However, regardless of whether these differences resulted from losses or gains, it is clear that the changes in female song and plumage must have occurred quite rapidly.

Moreover, it is also clear from our study that the majority of recent evolutionary changes in these taxa and in many other icterids have occurred in females rather than males. In the Agelaius genus, for instance, females differ substantially between species in both their vocal behaviours and plumage colours, some being nearly indistinguishable from males (A. assimilis, Agelaius xanthomus and Agelaius humeralis) and others being strikingly different (A. phoeniceus and Agelaius tricolor), while all males produce similar buzzy songs and have black plumage with brightly coloured epaulets (Whittingham et al. 1992, 1996; Beedy & Hamilton 1999; Jaramillo & Burke 1999; Barker et al. 2008). A comparable pattern is seen in the orioles, in which differences between the sexes in song and plumage have also resulted from historical changes in females rather than males (figures 1 and 2; Hofmann et al. 2008; also see Irwin 1994). Our findings should therefore caution future studies of sexual selection against focusing too much on male traits. For example, the red-winged blackbird has long served as a model for investigating the function and evolution of male songs and colours (Searcy & Yasukawa 1995; Catchpole & Slater 2008). Yet it is likely that both traits have changed relatively little in males during the evolution of this species and that most differences between the sexes have resulted from selection on females.

Our evidence that singing by both sexes was the ancestral state for many temperate New World blackbirds is consistent with a recent phylogenetic study by Garamszegi et al. (2007), showing that female song was present in the ancestors of many European songbirds that now lack female song. That study did not, however, investigate ancestral breeding ranges (Garamszegi et al. 2007). Some authors have further suggested that song in both sexes was the ancestral state for all oscine passerines (Riebel et al. 2005), based on evidence for an Australasian origin of songbirds (Barker et al. 2002) combined with the prevalence of female song among Australian species today. These possibilities present interesting challenges for future researchers to identify the factors favouring loss of female song, and they further emphasize the importance of understanding, whether male-specific song in a species is primarily due to selection for complex song in males or selection against such songs in females.

The disagreement between parsimony and maximumlikelihood reconstructions in our study underscores the importance of not relying on any single method in ancestral reconstructions (Cunningham et al. 1998). Both methods supported similar patterns of character change in breeding latitude across the family and in female song among the orioles, but they largely disagreed about ancestral changes in the oropendolas and caciques and in the grackles and allies. Maximum-likelihood uses information about phylogenetic branch lengths while parsimony does not, and this difference presumably explains much of the disagreement in these results, since both clades include relatively short internodes in their early histories (figure 2). Using parsimony alone therefore could have provided potentially misleading conclusions about many of these ancestral states, especially because parsimony reconstructions imply no uncertainty about the states of these nodes. But maximum-likelihood reconstructions alone might also be misleading, since they suggest both low female song rates and tropical breeding ranges in many deep ancestral nodes, which contradict the strong overall pattern of high rates of female song in tropical breeders today. Both the concentrated changes test and Pagel's (1994) test for correlated trait evolution supported a clear relationship between female song rates and breeding latitude, showing that such comparative methods can work well even when ancestral reconstruction methods fail to agree. Indeed, large numbers of evolutionary changes on a tree, such as those characterizing female song rates and breeding latitudes in the Icteridae, can provide more power for statistical comparative methods while at the same time reducing our confidence in ancestral states (Oakley & Cunningham 2000).

Although we found a strong general relationship between female song rates and breeding latitude across the Icteridae, the factors favouring each loss or gain in female song were not always clear and indeed may have differed among lineages. For example, unlike the many taxa that appear to have lost female song with the evolution of temperate migration, both the oropendola and cacique clade and the cowbird (Molothrus) clade present cases in which female song was apparently lost in tropical resident taxa (figure 1). In the oropendolas and caciques, species that lack female song are also known to be polygynous and nest colonially (Feekes 1981, 1982; Trainer 1989; Webster 1994; Price & Lanyon 2004; Fraga & Kreft 2007), while taxa with frequent female song appear to form year-round territorial pairs (Robinson 1986; Skutch 1996; Jaramillo & Burke 1999). The cowbirds, on the other hand, are interspecific brood parasites, a trait that presumably evolved in the ancestor of the genus (Jaramillo & Burke 1999; Searcy et al. 1999). Thus, in both the oropendolas and caciques and the cowbirds, female song appears to have been lost with changes in breeding behaviours, possibly from monogamous ancestors in which both sexes shared territorial duties and parental investment. These exceptions to the relationship between female song rates and breeding latitude suggest that it is not latitude per se influencing changes in female song, but rather other aspects of life history that tend to be associated with temperate or tropical habitats (Slater & Mann 2004). Future, more detailed work on this family should include additional characters such as mating system, territorial behaviours and migration to investigate the strength of any associations with female song.

Studies of individual species will also be needed to better understand the selective mechanisms behind the evolutionary changes revealed here.

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APPENDIX A.

species	female song	latitude ^a	references
Agelaioides badius	infrequent	S temperate ^b	Hauber et al. (1999) and Jaramillo & Burke (1999
Agelaius assimilis	duet with male	tropical	Whittingham et al. (1992, 1997) and Jaramillo & Burke (1999)
Agelaius humeralis	duet with male	tropical	Whittingham et al. (1996) and Jaramillo & Burke (1999)
Agelaius phoeniceus	infrequent	N temperate	Yasukawa & Searcy (1995) and Jaramillo & Burke (1999)
Agelaius tricolor	infrequent	N temperate	Beedy & Hamilton (1999) and Jaramillo & Burke (1999)
Agelaius xanthomus	similar to male	tropical	Jaramillo & Burke (1999)
Agelasticus thilius	rare or absent	S temperate	Jaramillo & Burke (1999)
Agelasticus xanthophthalmus	similar to male	tropical	Orians & Orians (2000)
Amblycercus holocericeus	duet with male	tropical	Ridgely & Tudor (1989), Howell & Webb (1995) and Jaramillo & Burke (1999)
Amblyramphus holosericeus	infrequent	S temperate ^b	Jaramillo & Burke (1999)
Cacicus cela	infrequent	tropical	Feekes (1981, 1982), Trainer (1989) and Jaramillo & Burke (1999)
Cacicus chrysopterus	duet with male	S temperate ^b	Stiles & Skutch (1989), Skutch (1996) and Jaramille & Burke (1999)
Cacicus haemorrhous	rare or absent	tropical	Feekes (1981) and Jaramillo & Burke (1999)
Cacicus solitarius	duet with male	tropical	Ridgely & Tudor (1989) and Jaramillo & Burke (1999)
Cacicus uropygialis	duet with male	tropical	Stiles & Skutch (1989), Skutch (1996) and Jaramille & Burke (1999)
Chrysomus icterocephalus	rare or absent	tropical	Wiley & Wiley (1980) and Jaramillo & Burke (1999
Chrysomus ruficapillus	rare or absent	S temperate ^b	Jaramillo & Burke (1999)
Curaeus forbesi	similar to male	tropical	Skutch (1996) and Jaramillo & Burke (1999)
Dives dives	duet with male	tropical	Orians (1983), Stiles & Skutch (1989) and Jaramille & Burke (1999)
Dives warszewiczi	duet with male	tropical	Orians (1983) and Jaramillo & Burke (1999)
Dolichonyx oryzivorus	rare or absent	N temperate	Martin & Gavin (1995) and Jaramillo & Burke (1999)
Euphagus carolinus	infrequent	N temperate	Avery (1995) and Jaramillo & Burke (1999)
Euphagus cyanocephalus	infrequent	N temperate	Jaramillo & Burke (1999) and Martin (2002)
Gnorimopsar chopi	similar to male	tropical ^b	Jaramillo & Burke (1999)
Gymnomystax mexicanus	similar to male	tropical	Skutch (1996) and Jaramillo & Burke (1999)
Gymnostinops bifasciatus	rare or absent	tropical	Jaramillo & Burke (1999) and Fraga & Kreft (2007
Gymnostinops guatimozinus	rare or absent	tropical	Jaramillo & Burke (1999)
Gymnostinops montezuma	rare or absent	tropical	Jaramillo & Burke (1999) and Price et al. (2006)
Icterus bullockii	similar to male	N temperate	Miller (1931) and Rising & Williams (1999)
Icterus chrysater	similar to male	tropical	Howell (1972) and Jaramillo & Burke (1999)
Icterus chrysocephalus	duet with male duet with male	tropical	Jaramillo & Burke (1999)
Icterus croconotus		tropical	Jaramillo & Burke (1999)
Icterus cucullatus	infrequent	N temperate	Pleasants & Albano (2001)
Icterus galbula	infrequent	N temperate	Beletsky (1982), Rising & Flood (1998) and Jaramillo & Burke (1999)
Icterus graduacauda	similar to male	tropical ^b	Flood (1990), Jaramillo & Burke (1999) and Flood et al. (2002)
Icterus icterus	similar to male	tropical	Skutch (1996) and Jaramillo & Burke (1999)
Icterus leucopteryx	similar to male	tropical	Scott (1893) and Jaramillo & Burke (1999)
Icterus mesomelas	duet with male	tropical	Stiles & Skutch (1989) and Jaramillo & Burke (1999)
Icterus parisorum	infrequent	N temperate	Scott (1885), Bent (1958) and Jaramillo & Burke (1999)
Icterus pectoralis	similar to male	tropical	Stiles & Skutch (1989), Skutch (1996) and Jaramille & Burke (1999)

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Appendix A. (Continued.)

species	female song	latitude ^a	references
Icterus pustulatus	similar to male	tropical ^b	Jaramillo & Burke (1999) and Price et al. (2008)
Icterus spurius	infrequent	N temperate	Bent (1958), Scharf & Kren (1996) and Jaramillo & Burke (1999)
Macroagelaius imthurni	similar to male	tropical	Jaramillo & Burke (1999)
Molothrus aeneus	rare or absent	N temperate ^b	Lowther (1995), Skutch (1996), Jaramillo & Burke (1999) and Warren (2002)
Molothrus ater	rare or absent	N temperate	Lowther (1993) and Jaramillo & Burke (1999)
Molothrus bonariensis	rare or absent	S temperate ^b	Skutch (1996), Hauber <i>et al.</i> (1999), Jaramillo & Burke (1999) and Lowther & Post (1999)
Molothrus oryzivorus	rare or absent	tropical	Stiles & Skutch (1989), Skutch (1996) and Jaramillo & Burke (1999)
Molothrus rufoaxillaris	rare or absent	S temperate	Ridgely & Tudor (1989), Skutch (1996), Hauber et al. (1999) and Jaramillo & Burke (1999)
Nesopsar nigerrimus	similar to male	tropical	Wiley & Cruz (1980), Skutch (1996) and Jaramillo & Burke (1999)
Psarocolius angustifrons	rare or absent	tropical	Jaramillo & Burke (1999) and Fraga & Kreft (2007)
Psarocolius atrovirens	rare or absent	tropical	Jaramillo & Burke (1999)
Psarocolius decumanus	rare or absent	tropical	Jaramillo & Burke (1999)
Psarocolius oseryi	rare or absent	tropical	Ridgely & Tudor (1989) and Jaramillo & Burke (1999)
Psarocolius viridis	rare or absent	tropical	Jaramillo & Burke (1999)
Psarocolius wagleri	rare or absent	tropical	Jaramillo & Burke (1999)
Quiscalus lugubris	rare or absent	tropical	Ridgely & Tudor (1989)
Quiscalus major	infrequent	N temperate	Post et al. (1996)
Quiscalus mexicanus	rare or absent	N temperate ^b	Skutch (1996), Jaramillo & Burke (1999) and Johnson & Peer (2001)
Quiscalus nicaraguensis	infrequent	tropical	Jaramillo & Burke (1999)
Quiscalus niger	infrequent	tropical	Jaramillo & Burke (1999)
Quiscalus quiscula	infrequent	N temperate	Wiley (1976) and Jaramillo & Burke (1999)
Sturnella magna	infrequent	N temperate ^b	Lanyon (1995) and Jaramillo & Burke (1999)
Sturnella neglecta	rare or absent	N temperate	Lanyon (1994) and Jaramillo & Burke (1999)
Xanthocephalus xanthocephalus	infrequent	N temperate	Twedt & Crawford (1995) and Jaramillo & Burke (1999)
Xanthopsar flavus	infrequent	S temperate	Jaramillo & Burke (1999) and Fraga (2005)

^a Scored based on whether more or less than 80 per cent of breeding range is above or below 23.5° latitude north (N) or south (S), based on range maps in Jaramillo & Burke (1999).

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^bSpecies in which less than 80 per cent of breeding range is included in either the tropical or a temperate zone; breeding ranges in these taxa were classified based on the locations of published observations of singing behaviour.

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