ANIMAL GENETICS

Widely Applicable PCR Markers for Sex Identification in Birds¹

M. N. Romanov^{*a*, *}, A. M. Betuel^{*b*}, L. G. Chemnick^{*c*}, O. A. Ryder^{*c*}, R. O. Kulibaba^{*d*}, O. V. Tereshchenko^{*e*}, W. S. Payne^{*f*}, Ph. C. Delekta^{*f*}, J. B. Dodgson^{*f*}, E. M. Tuttle^{*b*}, and R. A. Gonser^{*b*}

^aComparative and Functional Genomics Group, School of Biosciences, University of Kent, Canterbury, Kent, CT2 7NJ UK

^bThe Center for Genomic Advocacy, Department of Biology, Indiana State University, Terre Haute, IN 47809 USA

^cGenetics Division, San Diego Zoo Institute for Conservation Research, 15600 Escondido, CA 92027 USA

^dInstitute of Animal Science, National Academy of Agrarian Sciences of Ukraine, Kulinichi, Kharkov District, 62404 Ukraine

^eState Poultry Research Station, National Academy of Agrarian Sciences of Ukraine, Birky, Kharkiv region, 63421 Ukraine

^fDepartment of Microbiology and Molecular Genetics, Biomedical and Physical Sciences Bldg.,

Michigan State University, East Lansing, MI 48824 USA

*e-mail: m.romanov@kent.ac.uk

Received April 9, 2018; revised May 5, 2018; accepted May 29, 2018

Abstract—To aid in avian sex determination if birds are not sexually dimorphic and/or they are sexually immature, several molecular assays involving the polymerase chain reaction (PCR) have been developed. To test in a variety of domestic and wild avian species applicability of five sexing assays: previously described four assays based on nucleotide sequence differences between the Z and W copy of the chicken chromodomain-helicase-DNA-binding protein gene (*CHD1Z* and *CHD1W*), and a new sexing marker using the ubiquitin associated protein 2 (*UBAP2*) gene sequence. At least one molecular sexing marker was successful in 84 out of 88 examined species across 13 avian orders. These assays may be useful in breeding management of domestic and wild birds as well as in studies of avian ecology, population genetics, embryology and transgenesis.

Keywords: Aves, birds, DNA markers, PCR-based sexing **DOI:** 10.1134/S1022795419020121

INTRODUCTION

Molecular sex identification methods in domestic and wild birds are essential for species in which no or weak sexual dimorphism is evident either at hatch, maturity or *in ovo*. The usefulness of DNA-based sex determination has been demonstrated for evolutionary studies, ecological and conservation issues, and management of endangered species in the wild and captivity (e.g. [1-6]). Previously used sex-determination methods, involving traditional autosexing based on external traits, vent sexing and surgical gonad examination (e.g. [7]) and, later, sex-specific analyses of karvotypes and the amount of DNA per cell (e.g. [8– 10]) have been supplemented or replaced with genderspecific DNA fingerprints. Because in birds the homogametic sex is the male, with two Z chromosomes, and the heterogametic sex is the female, with one Z and one W chromosome, these techniques primarily employ repetitive (e.g. [11-14]), non-repetitive [15] or coding regions on the W chromosome that are absent or different from their homologs on the Z chromosome.

Cath

Polymerase chain reaction (PCR)-based markers provide rapid and inexpensive assays for avian Z/W sex determination. Most of these markers depend on the polymorphism between two conserved gene homologs, *CHD1Z* and *CHD1W* (e.g. [16–18]), both encoding a chromodomain-helicase-DNA-binding protein that plays an important role in gene regulation. The aim of this study was to examine the range of applicability of four existing *CHD1*-based sex markers [19–22] in poultry and wild avian species, and to expand the repertoire of molecular genetic markers that are applicable for determining gender in young and adult birds, embryos, and cell cultures.

MATERIALS AND METHODS

Experimental Birds and DNA Samples

In all, we examined 88 domestic and wild avian species across 13 orders (Tables 1–5) including: Galliformes (nine species), Anseriformes (two species), Cuculiformes (four species), Psittaciformes (29 species), Strigiformes (one species), Columbiformes (seven species), Gruiformes (two species), Ciconiiformes (three species), Falconiformes (seven species), Cathartiformes (one species), Charadriiformes (two

¹ The article is published in the original.

Species (by order)	P2/P3
Ciconiiformes	
African Openbill Stork (Anastomus lamelligerus)	+
Painted Stork (Mycteria leucocephala)	+
Columbiformes	
Coroneted Fruit-dove (Ptilinopus coronulatus trigeminus)	+
Western Orange-bellied Fruit-dove (P. iozonus humeralis)	+
Jambu Fruit-dove (P. jambu)	+
Sulawesi Superb Fruit-dove (P. superbus temminckii)	+
Pink-necked Green Pigeon (Treron vernans purpurea)	+
Falconiformes	
Harpy Eagle (Harpia harpyja)	+
Gruiformes	
Sandhill Crane (Grus canadensis)	+
Kagu (<i>Rhynochetos jubatus</i>)	+
Passeriformes	
San Clemente loggerhead shrike (Lanius ludovicianus mearnsi)	+
Po'ouli (<i>Melamprosops phaesoma</i>)	+
Procellariiformes	
Black-footed albatross (Phoebastria nigripes)	+
Psittaciformes	
Cockatoo (<i>Cacatua</i> sp.)	+

Table 1	Test regults main a the	Criffithe and Timeri	(1005) covin	a maina and in 1/	avian analias (auhan agias
таріе г.	Test results using the	CITERENTS AND THWAT	(1990) Sex II	19 Drimers in 14	avian species/subspecies

"+", sex dimorphism in band pattern.

species), Procellariiformes (one species), and Passeriformes (20 species). Domestic Chicken (Smoky line; [23]) and its ancestor, Red Jungle Fowl (UCD001 line; [24]) (Gallus gallus), Turkey (Meleagris gallopavo), Indian Peafowl (Pavo cristatus), Northern Bobwhite (Colinus virginianus), Common Quail (Coturnix coturnix) and Mallard (Anas platyrhynchos) samples were obtained from the Michigan State University poultry farm, East Lansing, MI, USA; Racing Homer pigeons (Columba livia) and House Wren (Troglodytes aedon) being also sampled for analysing at the Michigan State University. A wide range of birds was sampled at the San Diego Zoo and San Diego Zoo Safari Park, Escondido, CA, USA. Additional avian samples were obtained from the collections at Indiana State University as well as at the State Poultry Research Station, Birky, Ukraine, and Kharkiv Zoo, Kharkiv, Ukraine.

One mL or 200 μ L (in the case of small birds like the House Wren) whole blood samples of at least one male and/or one female of each species were collected, and DNA was phenol-chloroform extracted following standard protocols (e.g. [25]). Similar procedures were employed for DNA extraction from Domestic Chicken embryo tissues (spleen, gonad) and avian myeloblastosis virus-transformed cell lines. The latter were derived from six-day-old transgenic ADOL Line 0 chicks (J.B. Dodgson and W.S. Payne, unpublished data). Selected BAC clones were identified in a gridded genomic Red Jungle Fowl BAC library (TAMU library code 031-JF256-BI; [26]), and their DNAs were isolated as described elsewhere [27]. Blood samples of Ukrainian origin birds were obtained using DNA-Sorb-B (InterLabService, Moscow, Russian Federation) filter paper and were treated according to the manufacturer's protocol. DNA was eluted from the sorbent using TE buffer (50 μ L). A subset of samples from Indiana State University were extracted from tissue using the magnetic extraction DNA IQTM System (Promega, Madison, WI, USA). A single individual from certain species was used for testing if the specimen demonstrated sexually dimorphic plumage.

PCR Procedure and Primer Design

PCR was performed in 10, 15, or 25 μ L volumes including up to 50 ng genomic DNA, 40 to 300 nM of each primer, 1× *Taq* polymerase buffer (Life Technologies, Rockville, MD, USA), 1.5 mM MgCl₂, 200 μ M dNTP, and 0.5 or 1 unit *Taq* polymerase (Life Technologies, Rockville, MD, USA). The settings for the thermal cycler were the following: hot start at 94°C (2 min) followed by 40 cycles including denaturing at

Species (by order)	P8/P2
Anseriformes	
Mallard (Anas platyrhynchos)	+
Black Swan (Cygnus atratus)	+
Cathartiformes ^a	
California Condor (Gymnogyps californianus)	+
Charadriiformes	
European Herring Gull (Larus argentatus)	+
American Woodcock (Scolopax minor)	В
Ciconiiformes	
White Stork (Ciconia ciconia)	+
Columbiformes	
Mourning Dove (Zenaida macroura)	В
Cuculiformes	
Violet Turaco (Musophaga violacea)	+
Northern Chestnut-breasted Malkoha (Phaenicophaeus curvirostris	+
singularis)	
Red-billed Malkoha (Ph. javanicus)	+
Guinea Turaco (Tauraco persa)	+
Falconiformes	
Cinereous Vulture (Aegypius monachus)	+ ^b
Saker Falcon (Falco cherrug)	_
Bearded Vulture (Gypaetus barbatus)	B ^c
Griffon Vulture (Gyps fulvus)	+ ^b

Ta

+

+

+

+

+

+

+

+

A

А

Α

Α

+

+

A

А

White-tailed Eagle (Haliaeetus albicilla)

White Eared Pheasant (Crossoptilon crossoptilon)

Steller's Sea Eagle (H. pelagicus)

Domestic Chicken (Gallus gallus)

Guinea Fowl (*Numida meleagris*)

Common Myna (Acridotheres tristis)

Dark-eyed Junco (Junco hyemalis)

Po'ouli (Melamprosops phaesoma)

House Sparrow (Passer domesticus)

Northern Cardinal (Cardinalis cardinalis)

American Crow (Corvus brachyrhynchos)

Common Yellowthroat (Geothlypis trichas)

Northern Waterthrush (Parkesia noveboracensis)

San Clemente Loggerhead Shrike (Lanius ludovicianus mearnsi)

Turkey (*Meleagris gallopavo*)

Kagu (Rhynochetos jubatus)

Silver Pheasant (Lophura nycthemera)

Common Pheasant (Phasianus colchicus)

Galliformes

Gruiformes

Passeriformes

Table 2. (Contd.)

Species (by order)	P8/P2
Ovenbird (Seiurus aurocapillus)	Α
American Redstart (Setophaga ruticilla)	А
Eastern Bluebird (Sialia sialis)	Α
European Starling (Sturnus vulgaris)	В
Peruvian Blue-necked Tanager (Tangara cyanicollis cyanicollis)	+
Brown Thrasher (Toxostoma rufum)	В
American Robin (Turdus migratorius)	А
Hooded Warbler (Wilsonia citrina)	А
White-throated Sparrow (Zonotrichia albicollis)	+
White-crowned Sparrow (Z. leucophrys)	А
Psittaciformes	
Blue-fronted Amazon (Amazona aestiva)	+
Orange-winged Amazon (A. amazonica)	+
Hyacinth Macaw (Anodorhynchus hyacinthinus)	+
Red-winged Parrot (Aprosmictus erythropterus)	+
Blue-and-yellow Macaw (Ara ararauna)	+
Green-winged Macaw (A. chloroptera)	+
Scarlet Macaw (A. macao)	+
Chestnut-fronted Macaw (A. severus)	+
Orange-fronted Parakeet (Aratinga canicularis)	+
Jandaya Parakeet (A. jandaya)	+
Sun Parakeet (A. solstitialis)	+
Port Lincoln Parrot (Barnardius z. zonarius)	+
Sulphur-crested Cockatoo (Cacatua galerita)	+
Salmon-crested Cockatoo (C. moluccensis)	+
Spix's Macaw (Cyanopsitta spixii)	+
Red-fan Parrot (Deroptyus accipitrinus)	+
Red-shouldered Macaw (Diopsittaca nobilis)	+
Galah (Eolophus roseicapilla)	+
Budgerigar (Melopsittacus undulatus)	+
Monk Parakeet (Myiopsitta monachus)	+
Cockatiel (Nymphicus hollandicus)	+
Eastern Rosella (Platycercus eximius)	+
Red-fronted Parrot (Poicephalus gulielmi)	+
Senegal Parrot (P. senegalus)	+
Regent Parrot (Polytelis anthopeplus)	+
Palm Cockatoo (Probosciger aterrimus)	+
Red-breasted Parakeet (Psittacula alexandri)	+
African Grey Parrot (Psittacus erithacus)	+
Strigiformes	
Great Horned Owl (Bubo virginianus)	В

"+", sex dimorphism; "--", no sex dimorphism in band pattern; A, correctly identified sex of a single individual with known sexual phe-notype; B, produced positive bands for sex determination but no phenotypic confirmation; F, failed PCR. ^a Order status is debatable. California Condor has also been placed in Falconiformes as well as Ciconiiformes. ^b Sex differences were only obtained by *Bam*HI digestion of the PCR products.

^c Two female samples were only available, and Z- and W-specific bands were produced after both PCR and post-PCR BamHI digestion.

223

Species (by order)	2550F/2718R
Anseriformes	
Mallard (Anas platyrhynchos)	_
Columbiformes	
Pigeon (Columba livia)	+
Galliformes	
Common Quail (Coturnix coturnix)	+
Bobwhite Quail (Colinus virginianus)	+
Red Jungle Fowl (Gallus gallus)	+
Domestic Chicken (G. gallus)	+
Turkey (Meleagris gallopavo)	+
Indian Peafowl (Pavo cristatus)	+
Passeriformes	
House Wren (Troglodytes aedon)	_

Tabla 3	Test results using the	Fridalfsson and	Ellegren (10	00) ceving	nrimers in	nine avian	species
Table 5.	Test results using the	i muonsson anu	Ellegich (17	77 Sexing	primers in	inne avian	species

"+", sex dimorphism; "-", no sex dimorphism in band pattern.

 $94^{\circ}C$ (30 s), primer annealing at an appropriate temperature (1 min), elongation at $72^{\circ}C$ (2 min), and final extension at $72^{\circ}C$ (10 min). All PCR assays were optimised and replicated to maximise reproducibility. As PCR enhancers in optimisation and experimental trials, tetramethyl ammonium chloride (TMAC) and bovine serum albumin (BSA) were exploited in several cases.

The following Domestic Chicken *CHD1*-based sexing PCR primers (Fig. 1) were tested: (1) P2 [5'-TCTGCATCGCTAAATCCTTT-3'] and P3 [5'-AGATATTCTGGATCTGATAGTGA-3'] [19] at annealing temperature 54°C using post-PCR digestion with restriction enzymes (Table 1); (2) P8 [5'-CTCCCAAGGATGAGRAAYTG-3'] and P2 [5'-TCTGCATCGCTAAATCCTTT-3'] [20] using annealing temperature 48°C (Table 2); (3) 1237L [5'-GAGAAACTGTGCAAAACAG-3'] and 1272H [5'-TCCAGAATATCTTCTGCTCC-3'] [21] using annealing temperature 56°C and 1 mM TMAC as a PCR enhancer (Table 3); and (4) 2550F [5'-GTTACT-GATTCGTCTACGAGA-3'] and 2718R [5'-ATT-GAAATGATCCAGTGCTTG-3'] [22], with the annealing temperature being 52°C (Table 4).

To seek new sexing markers, a Turkey *AD012W* cDNA sequence (NCBI GenBank Accession no. AY188758) was used that is homologous to the human ubiquitin associated protein 2 (*UBAP2*) gene (NM_018449) and, by analogy with the Domestic Chicken *UBAP2* genes (NM_001277104, NM_001277105), has both Z and W chromosome copies. Using the program PrimerSelect (computer software package Lasergene99, DNASTAR, Madison, WI, USA), the following primers were designed: 166F [5'-GGTGTACCGCCCTTGTTG-3'] and 279R [5'-CATTGGCAGCCTTGGATTGAA-3'] within the putative exon 1, and 815F [5'-CCTGATAT-CAGTGGCTCTGTCTA-3'] and 906R [5'-GAGG-

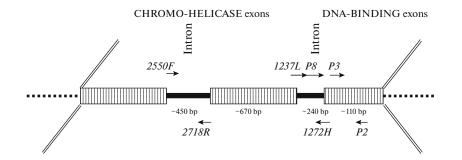


Fig. 1. Location of the *CHD1* gene-based sexing PCR primers (not on an exact scale): 2550F/2718R (Fridolfsson and Ellegren, 1999); Kahn 1237L/1272H (Kahn et al., 1998); P8/P2 (Griffiths et al., 1998); and P2/P3 (Griffiths and Tiwari, 1995). Exact positions of the primers on the chicken Z chromosome are: 2550F, 51,125,443–51,125,463; 2718R, 51,126,015–51,126,035; P8, 51,129,765–51,129,784; 1237L, 51,129,776–51,129,794; 1272H, 51,129,992–51,130,011; P3, 51,130,000–51,130,022; and P2, 51,130,090–51,130,109.

Table 4.	Test results using the K	ahn et al. (1998)) sexing primers	s in 33 avian	species/subspecies
Table 4.	Tool Toound adding the IN	ann et an (1990)	, sexing primere	m 55 umun	species/subspecies

Species (by order)	1237L/1272H
Anseriformes	
Mallard (Anas platyrhynchos)	_
Cathartiformes ^a	
California Condor (Gymnogyps californianus)	+
Charadriiformes	
American Woodcock (Scolopax minor) ^b	В
Columbiformes	
Pigeon (Columba livia)	F
Mourning Dove (Zenaida macroura) ^b	В
Galliformes	
Common Quail (Coturnix coturnix)	—
Bobwhite Quail (Colinus virginianus)	—
Red Jungle Fowl (Gallus gallus)	+
Domestic Chicken (G. gallus)	+
Turkey (Meleagris gallopavo)	+
Indian Peafowl (<i>Pavo cristatus</i>)	+
Gruiformes	
Kagu (<i>Rhynochetos jubatus</i>) Passeriformes	+
	А
Northern Cardinal (<i>Cardinalis cardinalis</i>) ^b	
American Crow (<i>Corvus brachyrhynchos</i>) ^b	A
Common Yellowthroat (Geothlypis trichas) ^b	A
Dark-eyed Junco (Junco hyemalis) ^b	A
San Clemente Loggerhead Shrike (Lanius ludovicianus mearnsi)	+
Po'ouli (Melamprosops phaesoma)	+
Northern Waterthrush (Parkesia noveboracensis) ^b	A
House Sparrow (Passer domesticus) ^b	А
Ovenbird (Seiurus aurocapillus) ^b	А
Black-throated Blue Warbler (Setophaga caerulescens) ^b	F
American Redstart (S. ruticilla) ^b	А
Eastern Bluebird (Sialia sialis) ^b	+
European Starling (<i>Sturnus vulgaris</i>) ^b	В
Peruvian Blue-necked Tanager (<i>Tangara c. cyanicollis</i>)	+
Brown Thrasher (<i>Toxostoma rufum</i>) ^b	В
House Wren (<i>Troglodytes aedon</i>)	F
American Robin (<i>Turdus migratorius</i>) ^b	А
Hooded Warbler (<i>Wilsonia citrina</i>) ^b	А
White-throated Sparrow (<i>Zonotrichia albicollis</i>) ^b	+
	A
White-crowned Sparrow (<i>Z. leucophrys</i>) ^b Strigiformes	2 1
-	В
Great Horned Owl (Bubo virginianus) ^b	ע

"+", sex dimorphism; "-", no sex dimorphism in band pattern; A, correctly identified sex of a single individual with known sexual phenotype; B, produced positive bands for sex determination but no phenotypic confirmation; F, failed PCR.

^a Order status is debatable. California Condor has also been placed in Falconiformes as well as Ciconiiformes.

^b The modified primers 1237L/1272Hzal were also used in this study.

GCATGCTGAAAGGAGGTG-3'] within the putative exon 3 (Table 5).

PCR products were analysed by running on 1 to 5% agarose gels in $1 \times$ TAE or TBE buffer, typically at

5 V/cm for 2 h, and visualizing with ethidium bromide under UV light. Alternatively, amplicons were analysed using capillary electrophoresis in an ABI PRISM 310[®] automated sequencer and analysed with the

Species (by order)	166F/279R		
Anseriformes			
Mallard (Anas platyrhynchos)	_		
Columbiformes			
Pigeon (Columba livia)	_		
Galliformes			
Common Quail (Coturnix coturnix)	_		
Bobwhite Quail (Colinus virginianus)	_		
Red Jungle Fowl (Gallus gallus)	+		
Domestic Chicken (G. gallus)	+		
Turkey (Meleagris gallopavo)	+		
Indian Peafowl (Pavo cristatus)	+		
Passeriformes			
House Wren (Troglodytes aedon)	_		

Table 5. Test results using the sexing primers based on UBAP2Z vs. UBAP2W sequence differences in nine avian species

"+", sex dimorphism; "-", no sex dimorphism in band pattern.

Genotyper® program (Applied Biosystems, Foster City, CA, USA).

CHD1 Sequencing

CHD1 PCR fragments were produced with 1237L and 1272H primers [21] using DNA templates from the Red Jungle Fowl, Turkey, Indian Peafowl, Northern Bobwhite, Common Quail and Mallard. The corresponding Z and W chromosome bands were excised from the gel, and DNA was extracted using QIAquick Gel Extraction Kit with microcentrifuge spin columns or QIAEX II Gel Extraction Kit using sodium iodide and silica-gel particles (QIAGEN, Valencia, CA, USA). All subcloning was done in the pCR®2.1-TOPO vector using the TOPO[™] TA Cloning[®] Kit (Invitrogen, Carlsbad, CA, USA). Sequencing was performed on an ABI377 sequencer (PerkinElmer, Boston, MA, USA) using cycle sequencing reactions with dye labelled M13 universal or reverse primers. The nucleotide sequence data reported in this paper have been submitted to the NCBI GenBank nucleotide sequence database and have been assigned the Accession no. AY775803.

Sequence Analysis

Local sequence alignments were performed using the Basic Local Alignment Search Tool (BLAST; [28]). Multiple alignments of avian *CHD1Z* and *CHD1W* sequences were done using Windows 32 MegAlign (DNASTAR, Madison, WI, USA). Alignments of sequences were performed using the CLUSTAL V and CLUSTAL W methods (as described in [29] and [30]), and were deposited in the EMBL-Align database [31] under accession numbers

ALIGN_000994-ALIGN_000996 ALIGN_000998-ALIGN_001000.

and

RESULTS

CHD1: P2/P3 Primers

We tested the original *CHD1*-based primer set [19] for sex identification based on the PCR-RFLP technique, i.e., PCR amplification coupled with post-PCR restriction enzyme digestion (Table 1). In all 14 species studied (belonging to seven avian orders), the sex dimorphism in the post-PCR restriction pattern was observed.

CHD1: P8/P2 Primers

We first validated this *CHD1*-based sexing marker on male and female pairs of four bird species: Common Pheasant (*Phasianus colchicus*), Budgerigar (*Melopsittacus undulates*), Red-fan Parrot (*Deroptyus accipitrinus*), and European Herring Gull (*Larus argentatus*) (Fig. 2). As expected, the male samples produced one amplicon, and the females two.

Furthermore, we confirmed the applicability of this test on 24 other birds of known sex including Domestic Chicken (Fig. 2a), Turkey, Guinea Fowl (Numida meleagris), Silver Pheasant (Lophura nycthemera) (Fig. 2a), African Grey Parrot (Psittacus erithacus) (Fig. 2b), White Stork (Ciconia ciconia) (Fig. 2a), American Woodcock (Scolopax minor), Great Horned Owl (Bubo virginianus), Mourning Dove (Zenaida macroura), and 15 passeriforms: Northern Cardinal (Cardinalis cardinalis), American Crow (Corvus brachyrhynchos), Common Yellowthroat (Geothlypis trichas), Dark-eyed Junco (Junco hyemalis), Northern Waterthrush (Parkesia

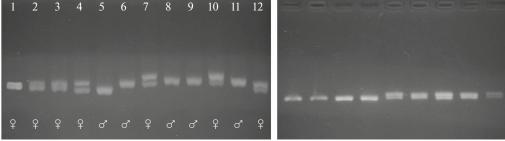


Fig. 2. PCR sexing assays using the P8/P2 primers: (a) Lanes: 1, Domestic Chicken female; 2, Indian Peafowl female; 3, Indian Peafowl female; 4, Common Pheasant female; 5, Common Pheasant male; 6, European Herring Gull male; 7, European Herring Gull female; 8, 9, White Stork males; 10, Budgerigar female; 11, Budgerigar male; 12, Silver Pheasant female. (b) Lanes: 1–4, African Grey Parrot males; 5, 7, 9, African Grey Parrot females; 6, 8, Red-fan Parrot females.

noveboracensis), House Sparrow (Passer domesticus), Ovenbird (Seiurus aurocapilla), American Redstart (Setophaga ruticilla), Eastern Bluebird (Sialia sialis), European Starling (Sturnus vulgaris), Brown Thrasher (Toxostoma rufum), American Robin (Turdus migratorius), Hooded Warbler (Setophaga citrina), Whitethroated Sparrow (Zonotrichia albicollis), and Whitecrowned Sparrow (Zonotrichia leucophrys) (data not shown).

(a)

To expand the range of the assay further, we tested a wider range of birds using DNA samples available at the San Diego Zoo, San Diego Zoo Safari Park, and Indiana State University. The results of molecular sexing tests in a total of 72 avian species/subspecies are summarised in Table 2. In all these trials, there was a significant variation in PCR product length depending on the species, although a common pattern of sex differences was observed in most cases, with males

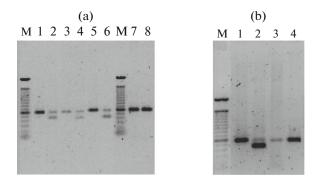


Fig. 3. PCR sexing assays using the 2550F/2718R primers (Fridolfsson and Ellegren 1999) and (a) 1.5 vs. (b) 3 mM MgCl₂ concentration: (a) Lanes: 1, Indian Peafowl male; 2, Indian Peafowl female; 3, Northern Bobwhite 1 male; 4, Northern Bobwhite female; 5, Common Quail male; 6, Common Quail female; 7, Domestic Chicken BAC (b) Lanes: 1, Turkey male; 2, Turkey female; 3, Mallard male; 4, Mallard female. M, 100-bp DNA ladder.

RUSSIAN JOURNAL OF GENETICS Vol. 55 2019 No 2

showing one PCR band and females showing two PCR bands. We were unable to identify sex-specific bands in the Saker Falcon (Falco cherrug), Whitetailed Eagle (Haliaeetus albicilla), and Steller's Sea Eagle (Haliaeetus pelagicus). There were also no sex differences in the Cinereous Vulture (Aegypius monachus) and Griffon Vulture (Gvps fulvus), so we additionally examined if distinguishing fragments could be obtained after post-PCR digestion with BamHI. The gender of two Bearded Vulture (Gypaetus barbatus) females was confirmed using both P8/P2 primers and post-PCR digestion with BamHI.

CHD1: 2550F/2718R Primers

When using primers developed by Fridolfsson and Ellegren [22], males of all galliform species studied displayed one 600 bp, Z-specific fragment, whereas females showed both CHD1Z (600 bp) and CHD1W (450-bp) fragments (Fig. 3, S4). Both Mallard sexes vielded a single product of 600 bp (Fig. 3b). Two CHD1-specific Red Jungle Fowl BACs. JB057N16 and JB059N09, also produced a Z copy fragment, thus confirming that they harboured the CHD1Z gene (Fig. 3a). Sexes were also distinguishable in the domestic pigeon using these primers (Fig. 4a) and generating a W fragment of similar size to those of the other birds, but a Z fragment was slightly larger (about 650 bp). However, this primer set failed to identify sexspecific bands in the House Wren (Fig. 4b; data shown for a female sample only). The test data for this sexing marker are summarised in Table 3.

CHD1: 1237L/1272H Primers

Using primers designed by Kahn et al. [21], three galliform species, Red Jungle Fowl/Domestic Chicken, Turkey and Indian Peafowl, displayed one Z-specific fragment (240 bp) in males and two fragments (240 and 265 bp) in females. Only a single frag-

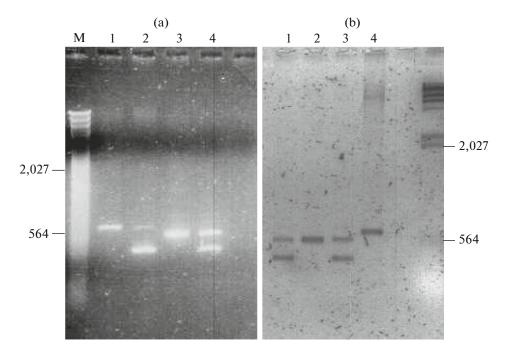


Fig. 4. PCR sexing assays using the 2550F/2718R primers (Fridolfsson and Ellegren 1999): (a) Lanes: 1, Pigeon male; 2, Pigeon female; 3, Red Jungle Fowl male; 4, Red Jungle Fowl female. M, 100-bp DNA ladder. (b) Lanes: 1, Red Jungle Fowl female; 2, Domestic Chicken Smoky line male; 3, Domestic Chicken Smoky line female; 4, House Wren female. M, λ /*Hin*dIII fragments.

ment was obtained, irrespective of sex, in Northern Bobwhite (250 bp), Common Quail (276 bp) and Mallard (240 bp) (data not shown). The Kahn et al. [21] primer amplification failed in Domestic Pigeon and House Wren samples of either sex.

A *CHD1* fragment amplified with Kahn et al. [21] primers was cloned and sequenced from the Common Quail only. However, its sequence (AY775803) was identical in both the male and female.

Based on the White-throated Sparrow BAC CH264-158I02 sequence (NCBI GenBank Accession no. AC237007), a modified reverse primer, 1272Hzal, was designed: [5'-TCCAGAGTATCTTCTACTCC-3']. It differed from the original reverse primer 1272 at the two specified nucleotides, and was utilised to improve the sexing assay for the White-throated Sparrow, especially in cases when DNA was recovered from old/difficult samples. We assessed both the original and modified reverse primer on a variety of Whitethroated Sparrow male and female samples as well as individual samples of 17 other species of known sex from four orders. The Northern Cardinal, Eastern Bluebird, and American Redstart specimens all demonstrated plumage characteristics that allowed for sex determination prior to genetic analysis. Our DNA results agreed with the dimorphic characteristics observed. Overall, this sexing marker efficiently worked in 30 out of 33 species tested (Table 4). Our data showed that the modified primer set worked perfectly in amplifying CHD1 fragments in the Whitethroated Sparrow and other passerines, even using old/difficult samples.

UBAP2: 166F/279R Primers

Based on the turkey UBAP2 cDNA sequence (AY188758), a new primer pair, 166F/279R, was designed that enabled molecular sexing in three galliform birds: Red Jungle Fowl, Turkey and Indian Peafowl (Fig. 5a; Table 4). A single amplified fragment in the males (Z copy) was \sim 550 bp long, whereas in the females there were two bands: \sim 550 bp (Z copy) and \sim 1000 bp (W copy). Similar fragments were obtained in sexing the Domestic Chicken embryo samples (spleen and gonad tissues) and cell lines (data not shown). On the other hand, there was a single amplification product in the Northern Bobwhite (1000 bp), Common Quail (550 bp) and Mallard (1000 bp) males and females (Fig. 5a). A single 1000-bp band was also obtained in both male and female Domestic Pigeon using the UBAP2 primers (Fig. 5b).

In the House Wren, the *UBAP2* primers generated an amplicon similar in the length to the *UBAP2W* gene fragment (\sim 1000 bp), regardless of sex (data not shown).

DISCUSSION

Most currently used molecular techniques for sexing birds involve differential PCR amplification in males and females of the chromodomain-helicase(a) M 1 2 3 4 5 6 7 8 9 10 11 12 M

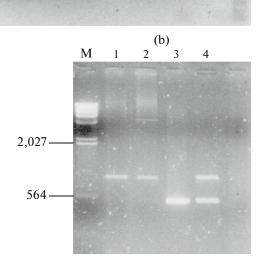


Fig. 5. PCR sexing assays using the *UBAP2*-based primers: (a) Lanes: 1, Red Jungle Fowl male; 2, Red Jungle Fowl female; 3, Turkey male; 4, Turkey female; 5, Indian Peafowl male; 6, Indian Peafowl female; 7, Northern Bobwhite male; 8, Northern Bobwhite female; 9, Common Quail male; 10, Common Quail female; 11, Mallard male (failed); 12, Mallard female; M, 100-bp DNA ladder. (b) Lanes: 1, Pigeon male; 2, Pigeon female; 3, Red Jungle Fowl male; 4, Red Jungle Fowl female. M, $\lambda/Hind III$ fragments.

DNA-binding gene with two sex chromosome-specific copies, CHD1Z and CHD1W, and we successfully tested four of these [19–22] across 13 avian orders. We also describe a novel sex-specific marker system based on the presence of Z and W copies of the UBAP2 gene [32–34]. This marker enabled us to accurately distinguish gender between males and females in three related galliform species: closelv Domestic Chicken/Red Jungle Fowl, Turkey, and Indian Peafowl. Its advantage compared to other methods of sexing birds is a pronounced size difference between the Z- (550 bp) and W-linked (1000 bp) amplified fragments. This provides a reliable method that does not require special gel electrophoresis conditions, as compared to the other tested markers with less obvious differences between the sex-specific bands, e.g. in the case of the Kahn et al. [21] primers producing a 25 bp difference. It also does not require application of the PCR-RFLP technique using post-PCR cleavage with restriction endonucleases, in order to distinguish between male and female PCR fragments of similar size, as was the case of the Griffiths and Tiwari [19] primer set and in a few species tested with [20] primers that required further *Bam*HI digestion.

Our data showed that the three widely domesticated galliform birds (Domestic Chicken/Red Jungle Fowl, Turkey, Indian Peafowl) can be sexed using any of the three tested DNA markers, and gender in two most common galliform poultry species, Domestic Chicken and Turkey, can be determined using four molecular markers (Tables 2–5). However, two quail species were successfully sexed in our hands with only the Fridolfsson and Ellegren [22] primers, and we were able to identify gender in the Mallard, the progenitor of domestic ducks, by means of the Griffiths et al. [20] primers alone.

In monomorphic avian species, in which both genders are similar, e.g. Domestic Pigeon and House Wren, development of molecular sexing markers would be beneficial. Although we succeeded in applying the Fridolfsson and Ellegren [22] primers for sex identification in the Domestic Pigeon, the methods tested failed in the House Wren. An effective approach would be to subclone and sequence the male and female amplified fragments that are of equal length but could potentially differ in nucleotide sequence at a certain restriction enzyme site. Indeed, Albrecht [35] reported a successful implementation of the Griffiths and Tiwari [19] marker coupled with the *Hae*III digestion that generated two smaller fragments of the W-linked fragments in the House Wren females.

Our experiments demonstrated that the choice of an appropriate, working molecular marker for sex identification in birds may depend on various factors including species tested, DNA sample/template quality, and primer/equipment availability. Use of enhancers/additives (e.g. TMAC) and a greater agarose gel concentration (up to 5%) could assist respectively for PCR optimisation and for a better amplicon band separation in cases when the sex chromosome-specific amplification fragments are very similar in size.

The molecular sexing assays described here and by other researchers can be useful in breeding management of domestic and wild galliforms and pigeons as well as in a variety of studies on other wild avian species including population genetics, population demography and structure, paternity verification, and conservation applications (e.g. [36]). They can also be effectively applied for identifying BAC clones that harbour sex-linked genes and markers in molecular genetics and genomics studies (e.g. physical mapping; [26], Romanov and Dodgson 2006) and for gene sequencing (as shown in this study for the Common Quail *CHD1* gene fragment). It is noteworthy that we efficiently employ molecular sex markers in the research projects focusing on the endangered California Condor and the White-throated Sparrow, a novel avian behavioural model (e.g. [38–40]). Also, the DNA tests for sex determination in chicken embryos and cell culture lines that were successful in our hands confirm utility of molecular sexing markers in embryology, transgenesis research, and search of *in ovo* sex identification techniques (e.g. [41]).

ACKNOWLEDGMENTS

The work was supported by grants from the United States Department of Agriculture/Cooperative State Research, Education, and Extension Service (99-35205-8566 and 2001-52100-11225) to J.B. Dodgson, and from the National Institutes of Health (R01GM084229) to E.M. Tuttle and R.A. Gonser. We are most grateful to Hans Ellegren (Uppsala University, Uppsala, Sweden) and Nate Kahn (University of Denver, Denver, CO, USA) for sharing aliquots of the sexing primers, 2550F/2718R and 1237L/1272H, respectively. We also thank Hans Ellegren for providing us with the UBAP2 sequence information, and Hans Cheng (USDA-ARS Avian Disease and Oncology Laboratory, East Lansing, MI, USA) for sequencing the common quail CHD1 fragments. Natalie Dubois (Michigan State University, East Lansing, MI, USA) is acknowledged for sharing the House Wren DNA samples, and Tanya Romanov (Michigan State University, East Lansing, MI, USA) and Sarah Ford (Indiana State University, Terre Haute, IN, USA) for technical assistance. We thank Olga Krestinina (Zelenodolsk, Russia) for graphical work.

COMPLIANCE WITH ETHICAL STANDARDS

Conflict of interests. The authors declare that they have no conflict of interest.

Statement on the welfare of animals. All applicable international, national, and/or institutional guidelines for the care and use of animals were followed.

REFERENCES

- Millar, C.D., Lambert, D.M., Anderson, S. and Halverson, J.L., Molecular sexing of the communally breeding pukeko: an important ecological tool, *Mol. Ecol.*, 1996, vol. 5, no. 2, pp. 289–293. https://onlinelibrary.wiley.com/doi/10.1046/j.1365-294X.1996.00076.x.
- Ellegren, H. and Fridolfsson, A.K., Male-driven evolution of DNA sequences in birds, *Nat. Genet.*, 1997, vol. 17, no. 2, pp. 182–184. https://doi.org/10.1038/ng1097-182.
- 3. Lessells, K., More mutations in males, *Nature*, 1997, vol. 390, no. 6657, pp. 236–237. https://doi.org/10.1038/36745.
- 4. Kahn, N.W. and Quinn, T.W., Male-driven evolution among Eoaves? A test of the replicative division hypothesis in a heterogametic female (ZW) system, J. *Mol. Evol.*, 1999, vol. 49, no. 6, pp. 750–759. https://doi.org/10.1007/PL00006597.

- Robertson, B.C., Millar, C.D., Minot, E.O., et al., Sexing the critically endangered kakapo *Strigops hab-roptilus, Emu*, 2000, vol. 100, no. 4, pp. 336–339. https://doi.org/10.1071/MU00056.
- Bermudez-Humaran, L.G., Garcia-Garcia, A., Leal-Garza, C.H., et al., Molecular sexing of monomorphic endangered *Ara* birds, *J. Exp. Zool.*, 2002, vol. 292, no. 7, pp. 677–680. https://doi.org/10.1002/jez.10070.
- Romanov, M.N. and Bondarenko, Y.V., Use of autosexing in waterfowl breeding and production, in *Proceedings of 10th European Symposium on Waterfowl*, Halle, 1995, pp. 473–476.
- Wang, N. and Shoffner, R.N., Trypsin G- and C-banding for interchange analysis and sex identification in the chicken, *Chromosoma*, 1974, vol. 47, no. 1, pp. 61–69. https://doi.org/10.1007/BF00326271.
- Nakamura, D., Tiersch, T.R., Douglass, M. and Chandler, R.W., Rapid identification of sex in birds by flow cytometry, *Cytogenet. Cell Genet.*, 1990, vol. 53, no. 4, pp. 201–205. https://doi.org/10.1159/000132930.
- De Vita, R., Cavallo, D., Eleuteri, P. and Dell'Omo, G., Evaluation of interspecific DNA content variations and sex identification in *Falconiformes* and *Strigiformes* by flow cytometric analysis, *Cytometry*, 1994, vol. 16, no. 4, pp. 346–350. https://doi.org/10.1002/cyto.990160409.
- Kagami, H., Nakamura, H. and Tomita, T., Sex identification in chickens by means of the presence of the W chromosome specific repetitive DNA units, *Jpn. Poult. Sci.*, 1990, vol. 27, no. 5, pp. 379–384. https://doi.org/10.2141/jpsa.27.379.
- Cassar, G., Mohammed, M., John, T.M., et al., Differentiating between parthenogenetic and "positive development" embryos in turkeys by molecular sexing, *Poult. Sci.*, 1998, vol. 77, no. 10, pp. 1463–1468. https://doi.org/10.1093/ps/77.10.1463.
- D'Costa, S. and Petitte, J.N., Sex identification of turkey embryos using a multiplex polymerase chain reaction, *Poult. Sci.*, 1998, vol. 77, no. 5, pp. 718–721. https://doi.org/10.1093/ps/77.5.718.
- Trefil, P., Bruno, M.M., Mikus, T. and Thoraval, P., Sexing of chicken feather follicle, blastodermal and blood cells, *Folia Biol.* (Praha), 1999, vol. 45, no. 6, pp. 253–256.
- Ogawa, A., Solovei, I., Hutchison, N., et al., Molecular characterization and cytological mapping of a nonrepetitive DNA sequence region from the W chromosome of chicken and its use as a universal probe for sexing Carinatae birds, *Chromosome Res.*, 1997, vol. 5, no. 2, pp. 93–101. https://doi.org/10.1023/A:1018461906913.
- Ellegren, H., First gene on the avian W chromosome (CHD) provides a tag for universal sexing of nonratite birds, *Proc. R. Soc. Lond. B Biol. Sci.*, 1996, vol. 263, no. 1377, pp. 1635–1641. https://doi.org/ 10.1098/rspb.1996.0239.
- Griffiths, R., Daan, S. and Dijkstra, C., Sex identification in birds using two CHD genes, *Proc. R. Soc. Lond. B Biol. Sci.*, 1996, vol. 263, no. 1374, pp. 1251–1256. https://doi.org/10.1098/rspb.1996.0184.
- Griffiths, R. and Korn, R.M., A *CHD1* gene is Z chromosome linked in the chicken *Gallus domesticus, Gene*, 1997, vol. 197, nos. 1–2, pp. 225–229. https://doi.org/10.1016/S0378-1119(97)00266-7.

- Griffiths, R. and Tiwari, B., Sex of the last wild Spix's macaw, *Nature*, 1995, vol. 375, no. 6531, p. 454. https://doi.org/10.1038/375454a0.
- 20. Griffiths, R., Double, M.C., Orr, K. and Dawson, R.J., A DNA test to sex most birds, *Mol. Ecol.*, 1998, vol. 7, no. 8, pp. 1071–1075. https://doi.org/10.1046/j.1365-294x.1998.00389.x.
- Kahn, N.W., St John, J. and Quinn, T.W., Chromosome-specific intron size differences in the avian CHD gene provide an efficient method for sex identification in birds, *Auk*, 1998, vol. 115, no. 4, pp. 1074–1078. https://doi.org/10.2307/4089527.
- Fridolfsson, A.K. and Ellegren, H., A simple and universal method for molecular sexing of non-ratite birds, *J. Avian Biol.*, 1999, vol. 30, no. 1, pp. 116–121. https://doi.org/10.2307/3677252.
- Kerje, S., Sharma, P., Gunnarsson, U., et al., The *Dominant white, Dun* and *Smoky* color variants in chicken are associated with insertion/deletion polymorphisms in the *PMEL17* gene, *Genetics*, 2004, vol. 168, no. 3, pp. 1507–1518. https://doi.org/10.1534/genetics.104.027995.
- Crittenden, L.B., Provencher, L., Santangelo, L., et al., Characterization of a Red Jungle Fowl by White Leghorn backcross reference population for molecular mapping of the chicken genome, *Poultry Sci.*, 1993, vol. 72, no. 2, pp. 334–348. https://doi.org/10.3382/ ps.0720334.
- Seutin, G., White, B.N. and Boag, P.T., Preservation of avian blood and tissue samples for DNA analyses, *Can. J. Zool.*, 1991, vol. 69, no. 1, pp. 82–90. https://doi.org/10.1139/z91-013.
- Lee, M.K., Ren, C.W., Yan, B., et al., Construction and characterization of three BAC libraries for analysis of the chicken genome, *Anim. Genet.*, 2003, vol. 34, no. 2, pp. 151–152. https://doi.org/10.1046/j.1365-2052.2003.00965_5.x.
- 27. https://web.archive.org/web/20100709231600/http:// hbz7.tamu.edu/homelinks/tool/bac_content.htm.
- Madden, T.L., Tatusov, R.L. and Zhang, J., Applications of network BLAST server, *Methods Enzymol.*, 1996, vol. 266, pp. 131–141.
- 29. Higgins, D.G. and Sharp, P.M., Fast and sensitive multiple sequence alignments on a microcomputer, *Comput. Appl. Biosci.*, 1989, vol. 5, no. 2, pp. 151–153. https://doi.org/10.1093/bioinformatics/5.2.151.
- Thompson, J.D., Higgins, D.G. and Gibson, T.J., CLUSTAL W: improving the sensitivity of progressive multiple sequence alignment through sequence weighting, position-specific gap penalties and weight matrix choice, *Nucleic Acids Res.*, 1994, vol. 22, no. 22, pp. 4673—4680. https://doi.org/10.1093/nar/22.22.4673.
- 31. Lombard, V., Camon, E.B., Parkinson, H.E., et al., EMBL-Align: a new public nucleotide and amino acid

multiple sequence alignment database, *Bioinformatics*, 2002, vol. 18, no. 5, pp. 763–764. https://doi.org/10.1093/bioinformatics/18.5.763.

- 32. Axelsson, E., Smith, N.G.C., Sundström, H., et al., Male-biased mutation rate and divergence in autosomal, Z-linked and W-linked introns of chicken and turkey, *Mol. Biol. Evol.*, 2004, vol. 21, no. 8, pp. 1538– 1547. https://doi.org/10.1093/molbev/msh157.
- Lawson Handley, L.J., Ceplitis, H. and Ellegren, H., Evolutionary strata on the chicken Z chromosome: implications for sex chromosome evolution, *Genetics*, 2004, vol. 167, no. 1, pp. 367–376. https://doi.org/10.1534/genetics.167.1.367.
- 34. Sazanov, A.A., Sazanova, A.L., Stekolnikova, V.A., et al., Chromosomal localization of the UBAP2Z and UBAP2W genes in chicken, Anim. Genet., 2006, vol. 37, no. 1, pp. 72–73. https://doi.org/10.1111/j.1365-2052.2005.01392.x.
- Albrecht, D.J., Sex ratio manipulation within broods of house wrens, *Troglodytes aedon, Anim. Behav.*, 2000, vol. 59, no. 6, pp. 1227–1234. https://doi.org/ 10.1006/anbe.1999.1420.
- Romanov, M.N., Kutnyuk, P.I. and Chernikov, V.F., Estimation of population structure and differentiation in black-headed gull by using genetic/oological parameters: 1. Analysis within an East-Ukrainian population, *J. Ornithol.*, 1994, vol. 135, no. 1, p. 261. https://doi.org/10.1007/BF02445773.
- Romanov, M.N. and Dodgson, J.B., Cross-species overgo hybridization and comparative physical mapping within avian genomes, *Anim. Genet.*, 2006, vol. 37, no. 4, pp. 397–399. https://doi.org/10.1111/j.1365-2052.2006.01463.x.
- Romanov, M.N., Koriabine, M., Nefedov, M., et al., Construction of a California condor BAC library and first-generation chicken—condor comparative physical map as an endangered species conservation genomics resource, *Genomics*, 2006, vol. 88, no. 6, pp. 711–718. https://doi.org/10.1016/j.ygeno.2006.06.005.
- Romanov, M.N., Tuttle, E.M., Houck, M.L., et al., The value of avian genomics to the conservation of wildlife, *BMC Genomics*, 2009, vol. 10, suppl. 2, p. S10. https://doi.org/10.1186/1471-2164-10-S2-S10.
- Romanov, M.N., Dodgson, J.B., Gonser, R.A. and Tuttle, E.M., Comparative BAC-based mapping in the white-throated sparrow, a novel behavioral model, using interspecies overgo hybridization, *BMC Res. Notes*, 2011, vol. 4, p. 211. https://doi.org/ 10.1186/1756-0500-4-211.
- 41. Narushin, V.G., Romanov, M.N. and Bogatyr, V.P., Method for preincubational prediction of embryo sex in chicken eggs, in *Proceedings of the 8th World Conference on Animal Production*, Seoul, 1998, vol. 1, pp. 832–833.