

A SURVEY OF AVAILABLE MOLECULAR MARKERS FOR VERTEBRATE SPECIES PRESENT IN COMUNIDAD DE MADRID

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ABSTRACT

In this paper I present a survey for molecular markers available for Vertebrate species inhabiting the Comunidad de Madrid, with the aim of being useful to researchers working on the characterization of animal biodiversity in this area.

Key words: Molecular markers, Vertebrate, Madrid.

RESUMEN

Estudio de los marcadores moleculares disponibles para las especies de vertebrados presentes en la Comunidad de Madrid

En este trabajo presento los resultados de una búsqueda en bases de datos de marcadores moleculares para especies de Vertebrados presentes en la Comunidad de Madrid, en la esperanza de que este trabajo pueda ser de utilidad para aquellos investigadores involucrados en la caracterización de biodiversidad animal en esta Comunidad.

Palabras clave: Marcadores moleculares, Vertebrados, Madrid.

Introduction

Today, molecular markers are a valuable tool for taxonomic and conservation studies allowing serious inferences in phylogeography, genetic structure of populations, conservation status, and so on (Bruford & Wayne, 1993; Tautz *et al.*, 2003; Avise, 2004).

For this reason, I consider that a survey of available molecular markers for Vertebrate species inhabiting in Comunidad de Madrid in sequence databases could be interesting as a research tool for biodiversity present in this area, giving researchers an overview to molecular knowledge for the species under scrutiny, and contributing to define gaps that can be filled subsequently.

The present survey is realised with this in mind, searching in sequence databases for mitochondrial, nuclear, and microsatellite (frequently repeated, highly polymorphic short sequences) sequences for vertebrate species or genera present in Comunidad de Madrid.

This approach, however, can have relative value only, since new sequences in sequence databases become available almost every day, and could make data shown in this work rapidly obsolete. In this sense, I believe that incorporating these data into a new database in progress in the Museo Nacional de Ciencias Naturales, and their continued review, can increase the value of the work.

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Table 1.— Sequences in databases for fish present in Comunidad de Madrid.

Tabla 1.— Secuencias en las bases de datos para peces presentes en la Comunidad de Madrid.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Ameiurus melas</i> / <i>Ictalurus melas</i>	13	12	1	0	85 sequences to the Genus (55753 sequences for <i>Ictalurus</i>) Many microsatellites
<i>Barbus bocagei</i>	28	25	3	0	733 sequences to the Genus 22 microsatellites
<i>Barbus comizo</i>	14	14	0	0	idem
<i>Carassius auratus</i>	745	yes	yes	21	790 sequences to the genus
<i>Chondrostoma arcasii</i> / <i>Rutilus arcasii</i>	29	29	0	0	185 sequences to the genus (55 for <i>Rutilus</i>) 5 microsatellites for <i>R. rutilus</i>
<i>Chondrostoma lemmingii</i> / <i>Rutilus lemmingii</i>	10	10	0	0	idem
<i>Chondrostoma polylepis</i>	4	4	0	0	idem
<i>Cobitis calderoni</i>	12	12	0	0	277 sequences to the Genus
<i>Cobitis paludica</i>	59	59	0	0	idem
<i>Cyprinus carpio</i>	11616	yes	yes	13	15 microsatellites for <i>C. auratus</i> are conserved
<i>Esox lucius</i>	108	42	40	26	171 sequences to the genus
<i>Gambusia holbrooki</i>	4	4	0	0	71 sequences to the Genus 7 microsatellites for <i>G. affinis</i> . 3 described for <i>G. holbrooki</i> are not in GenBank
<i>Gobio lozanoi</i> / <i>Gobio gobio</i>	58	44	14	0	67 sequences to the Genus
<i>Lepomis gibbosus</i>	11	8	3	0	215 sequences to the genus 9 microsatellites for <i>L. marginatus</i> , 1 for <i>L. macrochirus</i>
<i>Micropterus salmoides</i>	73	47	20	6	269 sequences to the genus. 11 microsatellites for <i>M. dolomieu</i>
<i>Oncorhynchus mykiss</i>	242135	yes	yes	967	246644 sequences to the Genus
<i>Silurus glanis</i>	21	3	2	16	42 sequences to the Genus
<i>Salmo trutta</i>	870	yes	yes	40	114040 sequences to the Genus
<i>Squalius alburnoides</i> / <i>Tropidophoxinellus alburnoides</i> / <i>Rutilus alburnoides</i>	67	67	0	0	149 sequences to the Genus (2 for <i>Tropidophoxinellus</i>) 6 microsatellites for <i>S. aradensis</i>
<i>Squalius pyrenaicus</i> / <i>Leuciscus pyrenaicus</i>	47	47	0	0	idem. (633 for <i>Leuciscus</i>) 5 microsatellites for <i>L. idus</i> and 5 for <i>R. rutilus</i> that are conser ved in <i>L. idus</i>
<i>Tinca tinca</i>	38	7	30	1	

Table 2.— Sequences in databases for Amphibia present in Comunidad de Madrid.

Tabla 2.— Secuencias en las bases de datos para anfibios presentes en la Comunidad de Madrid.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Alytes cisternasii</i>	11	5	0	6	105 sequences to the Genus 11 microsatellites for <i>A. muletensis</i>
<i>Alytes obstetricans</i>	57	51	6	0	idem. Genome mitochondrial completed
<i>Bufo bufo</i>	60	34	10	16	1772 sequences to the Genus
<i>Bufo calamita</i>	54	35	1	18	idem
<i>Discoglossus galganoi</i>	78	77	1	0	Genome mitochondrial completed 204 sequences to the Genus
<i>Discoglossus jeanneae</i>	72	71	1	0	idem
<i>Euproctus asper</i>	15	13	2	0	22 sequences to the Genus
<i>Hyla arborea</i>	25	4	4	17	1147 sequences to the genus
<i>Hyla meridionalis</i>	11	3	8	0	idem
<i>Pelobates cultripes</i>	22	15	7	0	37 sequences to the Genus
<i>Pelodytes punctatus</i>	21	17	4	0	30 sequences to the Genus
<i>Pleurodeles walt</i>	134	81	53	0	225 sequences to the Genus
<i>Rana iberica</i>	15	7	8	0	2022 sequences to the Genus Microsatellites
<i>Rana perezi</i>	12	9	3	0	idem
<i>Salamandra salamandra</i>	188	172	5	11	226 sequences to the Genus
<i>Triturus alpestris</i> ¹	10	2	1	7	427 sequences to the Genus 64 Microsatellites
<i>Triturus boscai</i> ²	6	6	0	0	idem
<i>Triturus marmoratus</i>	4	3	1	0	idem
<i>Triturus pygmaeus</i>	1	1	0	0	idem

(¹) Now *Mesotriton alpestris*. (²) Now *Lissotriton boscai*.

Methods

Each species present in Comunidad de Madrid has been searched for number of sequences, using the Taxonomy browser of the WEB page of NCBI (National Center for Biotechnology information USA) (<http://www.ncbi.nlm.nih.gov/Taxonomy/taxonomyhome.html>). This information has been edited, sorting them into mitochondrial and nuclear sequences and microsatellites when sequence number output is less than 200 sequences. Sequences available for the corresponding genera, with special focus in the presence of microsatellites, have also been analysed.

In cases with a sequence number output greater than 200, I search specifically for the presence of

microsatellite using the *Entrez* option with a query: "Genus name + microsatellite".

In the case of Fish I took the distribution show in the *Atlas y Libro Rojo de los Peces continentales de España* (Doadrio, 2002) as the basis of their presence in the region. For Amphibia, I followed Martínez-Solano & González Fernández (2003). In relation to Reptilia, I used the *Atlas y Libro Rojo de los Anfibios y Reptiles de España* (Pleguezuelos *et al.*, 2002). In the case of Aves, I split the species present in Madrid into sedentary and migrants on the basis of the *Anuario Ornitológico de Madrid 2002* (De la Puente Nilsson *et al.*, 2002), excluding species cited as accidental. For Mammals, I use the *Atlas de los mamíferos terrestres de España* (Palomo & Gisbert, 2002). For main synonymies I

Table 3.— Sequences in databases for Reptilia present in Comunidad de Madrid.

Tabla 3.— Secuencias en las bases de datos para reptiles presentes en la Comunidad de Madrid.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Acanthodactylus erithrurus</i>	41	41	0	0	87 sequences to the Genus
<i>Anguis fragilis</i>	8	3	5	0	
<i>Blanus cinereus</i>	3	0	3	0	5 sequences to the Genus
<i>Coluber hippocrepis</i> ¹	3	3	0	0	60 sequences to the Genus
<i>Coronella austriaca</i>	29	6	1	22	39 sequences to the Genus
<i>Coronella girondica</i>	10	9	1	0	22 microsatellites for <i>C. austriaca</i>
<i>Chalcides bedriagai</i>	0	0	0	0	139 sequences to the genus
<i>Chalcides striatus</i> / <i>Chalcides chalcides</i>	6	4	2	0	idem
<i>Chrysemys scripta</i>	8	0	8	0	95 sequences to the Genus
<i>Elaphe escalaris</i> ²	0	0	0	0	94 sequences to the Genus 11 microsatellites for <i>E. obsoleta</i>
<i>Emys orbicularis</i>	127	93	34	0	130 sequences to the Genus
<i>Iberolacerta cyreni</i> / <i>Lacerta monticola</i>	45	40	5	0	218 sequences to the Genus
<i>Lacerta lepida</i>	69	66	3	0	652 sequences to the Genus 9 microsatellites for <i>L. vivipara</i> and 12 for <i>L. viridis</i>
<i>Lacerta schreiberi</i>	36	32	4	0	idem
<i>Macroprotodon cucullatus</i>	25	23	2	0	109 sequences to the Genus
<i>Mauremys leprosa</i>	9	9	0	0	149 sequences to the Genus
<i>Mauremys caspica</i>	14	11	3	0	idem
<i>Malpolon monspessulanus</i>	13	10	3	0	16 sequences to the Genus
<i>Natrix maura</i>	21	20	1	0	155 sequences to the Genus 8 microsatellites for <i>N. tessellata</i>
<i>Natrix natrix</i>	70	68	2	0	idem
<i>Podarcis hispanica</i>	307	yes	yes	0	954 sequences to the Genus 9 microsatellites for <i>P. bocagei</i> , 9 for <i>P. muralis</i> and 4 for <i>P. erhardii</i>
<i>Podarcis muralis</i>	86	64	12	9	idem
<i>Psammodromus algirus</i>	46	42	4	0	48 sequences to the Genus
<i>Psammodromus hispanicus</i>	2	2	0	0	idem
<i>Tarentola mauritanica</i>	69	69	0	0	289 sequences to the Genus
<i>Vipera latastei</i>	2	2	0	0	352 sequences to the Genus 6 microsatellites for <i>V. berus</i>

⁽¹⁾ Now *Haemorrhois hippocrepis*. ⁽²⁾ Now *Rhinechis escalaris*.

grouped data for both entries (*Squalius pyrenai-cus/Leuciscus pyrenai-cus*). The data are current to November 16, 2005

Results

FISHES

Fishes species in Comunidad de Madrid are represented by a large number of molecular markers (Table 1). This result is logical if we consider that nine of twenty present species are exotic and with high economic value. Despite having a good number of sequences for *Cobitis*, no microsatellites has been described. Similarly, the absence of microsatellites for *Gobio* is remarkable. The taxonomic status of some species has been revised in recent years. In these cases, the search for the different synonymies shows a large number of molecular markers available. Additionally, a recent study (Larno *et al.*, 2005) characterizes new microsatellite markers for *Leuciscus* that are probably useful in other species (Barinova *et al.*, 2004).

AMPHIBIA

The 19 species present in Madrid are well represented in the GenBank (Table 2) except for *Triturus boscai*, *Triturus pygmaeus* and *Triturus marmoratus*, represented by 6, 1 and 4 sequences respectively and *Hyla meridionalis*, represented by 11 sequences. However, the Genus *Triturus* and *Hyla* are represented by a high number of sequences and among them they are numerous microsatellites that could be useful. All of the genera are well represented, with microsatellites described for *Alytes*, *Bufo*, *Hyla*, *Rana*, *Salamandra* and *Triturus*. In particular, microsatellites are described for *Alytes cisternasii*, *Bufo bufo*, *Bufo calamita*, *Hyla arborea*, *Salamandra salamandra* and *Triturus alpestris*.

Sequences are available from specimens sampled within the Comunidad de Madrid for *Alytes cisternasii*, *Alytes obstetricans*, *Discoglossus jeanneae* and *Salamandra salamandra* which may help in population studies.

REPTILIA

Most species present in Comunidad de Madrid are well represented by sequences in databases (Table 3), except *Chalcides bedriagai* and *Elaphe*

escalaris, and less so, *Blanus cinereus*, *Coluber hippocrepis*, *Psammodromus hispanicus* and *Vipera latastei*. However, they are many available sequences for the corresponding Genus except for *Blanus*.

With respect to microsatellites, a short number has been described for these species and genera. Only for *Coronella austriaca* and *Podarcis muralis* do we have microsatellites in GenBank, and at Genus level, microsatellites has been characterized for *Elaphe*, *Lacerta*, *Natrix*, *Podarcis* and *Vipera*. I believe that a supplementary effort in order to determine microsatellites for the Reptilia inhabiting the Comunidad de Madrid is necessary.

AVES

They are a large number of molecular markers for Aves, both sedentary (Table 4) and migrants (Table 5), that allow population, biogeographic and kinship studies. However, the number of markers is small for some genera. For this reason, it is necessary to implement the characterization of a larger number of molecular markers. For instance, there are no DNA markers for *Melanocorypha* among the sedentary group and other genera show a small number of markers (less than 10), including:

Sedentary:

<i>Aegyptius</i>	<i>Lullula</i>	<i>Pyrhocorax</i>
<i>Amandava</i>	<i>Miliaria</i>	<i>Remiz</i>
<i>Bubulcus</i>	<i>Oenanthe</i>	<i>Saxicola</i>
<i>Egretta</i>	<i>Petronia</i>	<i>Tachybaptus</i>
<i>Elanus</i>	<i>Phoenicurus</i>	<i>Tetrax</i>
<i>Garrulus</i>	<i>Ptyonoprogne</i>	<i>Upupa</i>

Migrants:

<i>Calandrella</i>	<i>Jinx</i>	<i>Pluvialis</i>
<i>Delichon</i>	<i>Lymnocyptes</i>	<i>Recurvirostra</i>
<i>Gelochelidon</i>	<i>Oenanthe</i>	<i>Saxicola</i>
<i>Himantopus</i>	<i>Philomachus</i>	<i>Scolopax</i>
<i>Ixobrychus</i>	<i>Phoenicurus</i>	<i>Tichodroma</i>

On the other hand, there are microsatellites available for the following genera: *Accipiter*, *Aegithalos*, *Alectoris*, *Anas*, *Aquila*, *Athene*, *Bubo*, *Buteo*, *Callipepla*, *Cettia*, *Columba*, *Corvus*, *Dendrocopos*, *Emberiza*, *Falco*, *Fringilla*, *Gyps*, *Hieraaetus*, *Larus*, *Loxia*, *Luscinia*, *Otis*, *Parus*, *Passer Phasianus*, *Phylloscopus*, *Pica*, *Podiceps*, *Prunella*, *Ptyonoprogne*, *Saxicola* and *Strix* within the sedentary group, and for genera: *Acrocephalus*, *Anas*,

Table 4.— Sequences in database for sedentary Aves present in Comunidad de Madrid.

Tabla 4.— Secuencias en bases de datos para aves sedentarias presentes en la Comunidad de Madrid.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Accipiter gentilis</i>	50	13	13	24	113 sequences to the Genus
<i>Accipiter nisus</i>	12	3	9	0	idem. 24 microsatellites for <i>A. gentilis</i>
<i>Actitis hypoleucos</i>	10	7	3	0	24 sequences to the genus
<i>Aegithalos caudatus</i>	16	10	5	1	21 sequences to the Genus
<i>Aegyptius monachus</i>	5	4	1	0	
<i>Alauda arvensis</i>	12	6	6	0	
<i>Alcedo atthis</i>	8	3	5	0	14 sequences to the Genus
<i>Alectoris rufa</i>	91	32	36	23	316 sequences to the Genus
<i>Amandava amandava</i>	2	2	0	0	7 sequences to the Genus
<i>Anas clypeata</i>	11	6	5	0	1879 sequences to the Genus.
<i>Anas platyrhynchos</i>	1111	yes	yes	More than 300	Microsatellites for <i>A. platyrhynchos</i>
<i>Anas strepera</i>	66	28	38	0	idem
<i>Anthus spinoletta</i>	3	1	2	0	32 sequences to the Genus
<i>Aquila adalberti</i>	22	4	0	18	208 sequences to the Genus
<i>Aquila chrysaetos</i>	30	18	12	0	8 microsatellites for <i>A. heliaca</i>
<i>Asio otus</i>	22	17	5	0	idem
<i>Athene noctua</i>	9	9	0	0	37 sequences to the Genus
<i>Aythya ferina</i>	1	1	0	0	17 sequences to the Genus.
<i>Bubo bubo</i>	48	28	13	7	7 microsatellites for <i>A. cunicularia</i>
<i>Bubulcus ibis</i>	4	3	1	0	18 sequences to the Genus
<i>Burhinus oedicephalus</i>	1	1	0	0	95 sequences to the genus
<i>Buteo buteo</i>	123	65	2	56	12 sequences to the Genus
<i>Callipepla californica</i>	19	11	1	7	Genome mitochondrial completed.
<i>Carduelis cannabina</i>	1	1	0	0	265 sequences to the Genus
<i>Carduelis carduelis</i>	6	5	1	0	35 sequences to the Genus. Exotic
<i>Carduelis chloris</i>	19	15	4	0	78 sequences to the Genus
<i>Certhia brachydactyla</i>	1	1	0	0	idem
<i>Cettia cetti</i>	6	6	0	0	idem
<i>Cinclus cinclus</i>	37	31	6	0	15 sequences to the Genus
<i>Circus aeruginosus</i>	9	5	4	0	31 sequences to the Genus.
<i>Circus cyaneus</i>	4	4	0	0	12 microsatellites for <i>C. diphone</i>
<i>Cisticola jundicis</i>	2	2	0	0	42 sequences to the Genus
<i>Coccothraustes coccothraustes</i>	6	3	3	0	20 sequences to the genus
<i>Columba livia</i>	318	yes	yes	18	idem
<i>Columba oenas</i>	1	1	0	0	11 sequences to the Genus
<i>Columba palumbus</i>	29	19	10	0	10sequences to the Genus
<i>Corvus corax</i>	282	yes	yes	1	459 sequences to the Genus
<i>Corvus corone</i>	17	8	9	0	idem. Microsatellites for <i>C. livia</i>
<i>Corvus monedula</i>	6	1	5	0	idem. 1 microsatellite for <i>C. corax</i>
<i>Cyanopica cooki</i>	8	8	0	0	idem
<i>Cyanopica cyanus</i>	65	62	3	0	73 sequences to the Genus
<i>Dendrocopos major</i>	276	yes	yes	0	idem
<i>Dendrocopos minor</i>	0	0	0	0	296 sequences to the Genus. 3 microsatellites for <i>D. leucopterus</i> and 6 for <i>D. leucotos</i>
<i>Egretta garzetta</i>	2	1	1	0	idem
<i>Elanus caeruleus</i>	0	0	0	0	8 sequences to the Genus
<i>Emberiza cioides</i>	4	4	0	0	3 sequences to the Genus
					117 sequences to the Genus. 26 microsatellites for <i>E. citrinella</i> and 7 for <i>E. schoeniclus</i>

Table 4.— Cont.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Emberiza cirius</i>	2	2	0	0	idem
<i>Erithacus rubecula</i>	79	68	11	0	
<i>Falco peregrinus</i>	180	108	43	29	Genome mitochondrial Completed. 328 sequences to the Genus. 28 microsatellites for <i>F. peregrinus</i> , 8 for <i>F. rusticolus</i> and 2 for <i>F. punctatus</i>
<i>Falco tinnunculus</i>	24	13	11	0	idem
<i>Fringilla coelebs</i>	87	70	16	1	121 sequences to the Genus 1 satellite
<i>Fulica atra</i>	3	2	1	0	11 sequences to the Genus
<i>Galerida cristata</i>	13	11	2	0	24 sequences to the Genus
<i>Galerida theklae</i>	9	7	2	0	idem
<i>Gallinula chloropus</i>	13	2	11	0	
<i>Garrulus glandarius</i>	1	1	0	0	4 sequences to the Genus
<i>Gyps fulvus</i>	19	7	2	10	39 sequences to the Genus
<i>Hieraaetus fasciatus</i>	52	12	5	35	91 sequences to the Genus
<i>Lanius excubitor</i>	16	13	3	0	132 sequences to the Genus
<i>Lanius meridionalis</i>	38	35	3	0	idem
<i>Larus ridibundus</i>	22	9	13	0	710 sequences to the genus. 7 microsatellites for <i>L. novaehollandiae</i>
<i>Loxia curvirostra</i>	43	39	4	0	58 sequences to the Genus. 7 microsatellites for <i>L. scotica</i>
<i>Lullula arborea</i>	2	2	0	0	
<i>Luscinia svecica</i>	302	yes	yes	2	310 sequences to the genus
<i>Melanocorypha calandra</i>	0	0	0	0	
<i>Miliaria calandra</i>	8	8	0	0	
<i>Milvus milvus</i>	46	46	0	0	137 sequences to the genus
<i>Monticola solitarius</i>	4	3	1	0	21 sequences to the genus
<i>Motacilla alba</i>	494	yes	yes	0	1629 sequences to the Genus
<i>Motacilla cinerea</i>	27	23	4	0	idem
<i>Myiopsitta monachus</i>	11	10	1	0	Exotic
<i>Oenanthe leucura</i>	0	0	0	0	2 sequences to the Genus
<i>Otis tarda</i>	220	yes	yes	8	
<i>Parus ater</i>	7	5	2	0	1283 sequences to the Genus. Microsatellites for <i>P. major</i> (23), <i>P. monticolus</i> (12) and <i>P. caeruleus</i> (12)
<i>Parus caeruleus</i>	268	yes	yes	12	idem
<i>Parus cristatus</i>	1	1	0	0	idem
<i>Parus major</i>	405	yes	yes	23	idem
<i>Passer domesticus</i>	76	20	46	10	129 sequences to the Genus
<i>Passer hispaniolensis</i>	4	4	0	0	idem. Microsatellites for <i>P. domesticus</i>
<i>Passer montanus</i>	29	18	11	0	idem
<i>Petronia petronia</i>	5	2	3	0	8 sequences to the Genus
<i>Phasianus colchicus</i>	67	13	53	1	78 sequences to the Genus. Exotic
<i>Phoenicurus ochrurus</i>	0	0	0	0	5 sequences to the Genus
<i>Phylloscopus collybita</i>	18	14	4	0	365 sequences to the Genus. 8 microsatellites for <i>P. occipitalis</i> , 2 for <i>P. trochilus</i> and 10 for <i>P. ijimae</i>
<i>Phylloscopus ibericus</i>	0	0	0	0	idem
<i>Pica pica</i>	50	39	7	4	60 sequences to the Genus
<i>Picus viridis</i>	3	2	1	0	14 sequences to the Genus
<i>Porphyrio porphyrio</i>	2	0	2	0	15 sequences to the Genus
<i>Podiceps cristatus</i>	2	0	2	0	36 sequences to the Genus. 7 microsatellites for <i>P. griseigena</i>
<i>Podiceps nigricollis</i>	1	1	0	0	idem
<i>Prunella collaris</i>	16	0	7	9	38 sequences to the Genus
<i>Prunella modularis</i>	11	4	7	0	idem. 9 microsatellites for <i>P. collaris</i>
<i>Pterocles alchata</i>	0	0	0	0	98 sequences to the Genus

Table 4.— End.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Pterocles orientalis</i>	1	0	1	0	idem
<i>Ptyonoprogne rupestris</i>	2	2	0	0	5 sequences to the Genus. 1 microsatellite for <i>P. fuligula</i>
<i>Pyrhacorax pyrrhacorax</i>	4	2	2	0	6 sequences to the Genus
<i>Rallus aquaticus</i>	3	3	0	0	12 sequences to the Genus
<i>Regulus ignicapillus</i>	4	4	0	0	54 sequences to the Genus
<i>Regulus regulus</i>	24	24	0	0	idem
<i>Remiz pendulinus</i>	7	2	5	0	
<i>Saxicola torquata</i>	2	2	0	0	8 sequences to the Genus. 1 microsatellite for <i>S. rubertra</i>
<i>Serinus citrinella</i>	1	1	0	0	73 sequences to the Genus
<i>Serinus serinus</i>	5	5	0	0	idem
<i>Sitta europaea</i>	287	yes	yes	0	392 sequences to the Genus
<i>Strix aluco</i>	328	yes	yes	0	417 sequences to the Genus. 7 microsatellites for <i>S. occidentalis</i>
<i>Sturnus unicolor</i>	0	0	0	0	32 sequences to the Genus
<i>Sylvia melanocephala</i>	5	5	0	0	54 sequences to the Genus
<i>Sylvia undata</i>	1	1	0	0	idem
<i>Tachybaptus ruficollis</i>	1	1	0	0	
<i>Tetrax tetrax</i>	6	4	2	0	
<i>Troglodytes troglodytes</i>	149	146	3	0	190 sequences to the Genus
<i>Turdus merula</i>	8	3	5	0	197 sequences to the Genus
<i>Turdus philomelos</i>	6	4	2	0	idem
<i>Turdus viscivorus</i>	1	0	1	0	idem
<i>Tyto alba</i>	34	24	10	0	59 sequences to the Genus
<i>Upupa epops</i>	7	4	3	0	
<i>Vanellus vanellus</i>	3	2	1	0	11 sequences to the Genus

Anser, Ardea, Calidris, Charadrius, Clamator, Corvus, Coturnix, Cuculus, Emberiza, Falco, Ficedula, Gallinago, Grus, Hirundo, Larus, Locustella Luscinia, Merops, Muscicapa, Otus, Phalacrocorax, Phylloscopus, Philomachus Phyrula, Riparia, Saxicola and Sterna among the migratory group.

It is necessary to remark the absence of microsatellites for *Turdus* and *Ciconia*, and the presence of only one microsatellite sequence for *Corvus*. The case of *Philomachus* is also interesting. In this genus, the nine sequences available are microsatellites.

MAMMALS

Mammals are well represented by sequences in databases except for the endangered species *Galemys pyrenaicus* with only four sequences deposited in GenBank.

In respect to the microsatellites, the number of sequences is large, except for *Barbastella, Elyomis,*

Genetta, Hypsugo, Neomys, Pipistrellus, Suncus and *Talpa*, with no microsatellites characterized.

Conclusions

The species and genera without molecular data deserve a particular effort for the characterization of molecular markers. In particular, I think that is necessary to invest in the obtention of more sequences from the endangered *Galemys pyrenaicus*, and to obtain sequences for underrepresented avian genera, such as *Elanus, Aegyptius, Melanocorypha, Lymnocyptes* and others.

On the other hand, it is also necessary to characterize microsatellites for genera with no sequences available, considering the current importance of these markers (Bruford & Wayne, 1993), and in particular, a major effort in the characterization of microsatellites in reptiles and fish should be undertaken.

Table 5.— Sequences in databases for migrant Aves present in Comunidad de Madrid.

Tabla 5.— Secuencias en bases de datos para aves migrantes presentes en la Comunidad de Madrid.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Acrocephalus arundinaceus</i>	71	6	53	12	294 sequences to the Genus. 67 microsatellites for <i>A. sechellensis</i> , 3 for <i>A. paludicola</i> and 1 for <i>A. orientalis</i>
<i>Acrocephalus schoenubaenus</i>	10	10	0	0	idem
<i>Acrocephalus scirpaceus</i>	12	12	0	0	idem
<i>Anas acuta</i>	11	8	3	0	1879 sequences to the genus. Microsatellites for <i>A. platyrhynchos</i>
<i>Anas crecca</i>	9	5	4	0	idem
<i>Anas penelope</i>	63	23	40	0	idem
<i>Anas querquedula</i>	5	5	0	0	idem
<i>Anser anser</i>	82	12	70	0	325 sequences to the Genus. 47 microsatellites para <i>A. cygnoides</i> and 2 for <i>A. albifrons</i>
<i>Anthus campestris</i>	1	1	0	0	32 sequences to the Genus
<i>Anthus pratensis</i>	9	6	3	0	idem
<i>Anthus trivialis</i>	5	2	3	0	idem
<i>Apus apus</i>	17	14	3	0	37 sequences to the Genus
<i>Apus pallidus</i>	0	0	0	0	idem
<i>Ardea cinerea</i>	5	2	3	0	76 sequences to the Genus. 60 microsatellites for <i>A. herodias</i>
<i>Ardea purpurea</i>	1	1	0	0	idem
<i>Asio flammeus</i>	15	12	3	0	37 sequences to the Genus
<i>Aythya fuligula</i>	2	1	1	0	18 sequences to the Genus
<i>Aythya nyroca</i>	1	1	0	0	idem
<i>Calandrella brachydactyla</i>	0	0	0	0	4 sequences to the Genus
<i>Calidris alpina</i>	43	41	2	0	341 sequences to the Genus. 3 microsatellites for <i>C. canutus</i>
<i>Calidris ferruginea</i>	2	2	0	0	idem
<i>Calidris minuta</i>	0	0	0	0	idem
<i>Caprimulgus europaeus</i>	3	2	1	0	15 sequences to the Genus
<i>Caprimulgus ruficollis</i>	0	0	0	0	idem
<i>Carduelis spinus</i>	3	3	0	0	78 sequences to the Genus
<i>Charadrius dubius</i>	1	0	1	0	183 sequences to the Genus. 66 microsatellites for <i>C. alexandrinus</i>
<i>Charadrius hiaticula</i>	0	0	0	0	idem
<i>Chlidonias hybridus</i>	3	3	0	0	10 sequence to the Genus
<i>Chlidonias niger</i>	4	4	0	0	idem
<i>Ciconia ciconia</i>	7	5	2	0	Genome mitochondrial completed. 74 sequences to the Genus
<i>Ciconia nigra</i>	15	10	5	0	idem
<i>Circaetus gallicus</i>	8	5	3	0	19 sequences to the Genus
<i>Circus pygargus</i>	0	0	0	0	20 sequences to the genus
<i>Clamator glandarius</i>	10	3	0	7	13 sequences to the Genus
<i>Coracias garrulus</i>	5	4	1	0	24 sequences to the Genus
<i>Corvus frugileus</i>	7	4	3	0	439 sequences to the Genus. 1 microsatellite for <i>C. corax</i> . Genome mitochondrial completed
<i>Coturnix coturnix</i>	294	yes	yes	0	648 sequences to the genus. 115 microsatellites for <i>C. japonica</i>
<i>Cuculus canorus</i>	45	32	6	7	63 sequences to the Genus
<i>Delichon urbica</i>	3	2	1	0	8 sequences to the Genus
<i>Emberiza citrinella</i>	39	6	7	26	117 sequences to the Genus. 26 microsatellites for <i>E. citrinella</i> and 7 for <i>E. schoeniclus</i>
<i>Emberiza hortelana</i>	0	0	0	0	idem
<i>Emberiza schoeniclus</i>	37	23	7	7	idem
<i>Falco columbarius</i>	14	4	10	0	328 sequences to the Genus. 28 microsatellites for <i>F. peregrinus</i> , 8 for <i>F. rusticolus</i> and 2 for <i>F. punctatus</i>

Table 5. Cont.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Falco eleanorae</i>	0	0	0	0	idem
<i>Falco naumanni</i>	1	1	0	0	idem
<i>Falco subbuteo</i>	3	3	0	0	idem
<i>Ficedula hypoleuca</i>	164	60	81	23	350 sequences to the genus. Other microsatellites
<i>Fringilla montifrigilla</i>	23	16	7	0	121 sequences to the Genus
<i>Gallinago gallinago</i>	7	3	4	0	118 sequences to the Genus. Microsatellites for <i>G. media</i>
<i>Gelochelidon nilotica</i>	1	1	0	0	
<i>Grus grus</i>	11	6	5	0	349 sequences to the Genus. 35 microsatellites for <i>G. americana</i> , 7 for <i>G. japonensis</i> , 1 for <i>G. paradisea</i> and 1 for <i>G. canadensis</i>
<i>Hieraaetus pennatus</i>	11	7	4	0	91 sequences to the Genus
<i>Himantopus himantopus</i>	1	0	1	0	4 sequences to the Genus.
<i>Hippolais polyglotta</i>	4	4	0	0	27 sequences to the Genus
<i>Hirundo daurica</i>	0	0	0	0	89 sequences to the Genus. Microsatellites for <i>H. rustica</i>
<i>Hirundo rustica</i>	45	14	15	16	idem
<i>Ixobrychus minutus</i>	0	0	0	0	5 sequences to the Genus
<i>Jynx torquilla</i>	4	3	1	0	
<i>Lanius collurio</i>	9	4	5	0	132 sequences to the Genus
<i>Lanius senator</i>	6	6	0	0	idem
<i>Larus cachimans</i>	57	57	0	0	710 sequences to the Genus. 7 microsatellites for <i>L. novaehollandiae</i>
<i>Larus fuscus</i>	72	62	10	0	idem
<i>Larus melanocephalus</i>	4	4	0	0	idem
<i>Limosa limosa</i>	4	4	0	0	27 sequences to the Genus
<i>Locustella luscinioides</i>	3	3	0	0	63 sequences to the Genus. 7 microsatellites for <i>L. pyeri</i>
<i>Locustella naevia</i>	7	7	0	0	idem
<i>Luscinia megarhynchos</i>	0	0	0	0	310 sequences to the Genus. 2 microsatellites for <i>L. svecica</i>
<i>Lymnocyptes minimus</i>	1	1	0	0	
<i>Merops apiaster</i>	20	0	0	20	30 sequences to the Genus
<i>Milvus migrans</i>	91	87	4	0	137 sequences to the Genus
<i>Monticola saxatilis</i>	0	0	0	0	21 sequences to the Genus
<i>Motacilla flava</i>	885	yes	yes	0	1629 sequences to the Genus
<i>Muscicapa striata</i>	23	5	15	3	37 sequences to the Genus
<i>Neophron percnopterus</i>	33	31	2	0	
<i>Numenius arquata</i>	2	2	0	0	13 sequences to the Genus
<i>Nycticorax nycticorax</i>	13	8	5	0	
<i>Oenanthe hispanica</i>	0	0	0	0	2 sequences to the Genus
<i>Oenanthe oenanthe</i>	1	1	0	0	idem
<i>Oriolus oriolus</i>	3	0	3	0	15 sequences to the Genus
<i>Otus scops</i>	10	6	4	0	118 sequences to the Genus. 18 microsatellites for <i>O. elegans</i>
<i>Pandion haliaetus</i>	16	8	8	0	
<i>Pernis apivorus</i>	7	7	0	0	38 sequences to the Genus
<i>Phalacrocorax carbo</i>	38	27	4	7	112 sequences to the Genus
<i>Philomachus pugnax</i>	9	0	0	9	
<i>Phoenicurus phoenicurus</i>	2	2	0	0	5 sequences to the Genus
<i>Phylloscopus bonelli</i>	5	5	0	0	365 sequences to the genus. 8 microsatellites for <i>P. occipitalis</i> , 2 for <i>P. trochilus</i> and 10 for <i>P. ijimae</i>
<i>Phylloscopus trochilus</i>	28	10	16	2	idem
<i>Platalea leucorodia</i>	0	0	0	0	267 sequences to the Genus
<i>Pluvialis apricaria</i>	0	0	0	0	7 sequences to the Genus
<i>Pyrrhula pyrrhula</i>	10	8	0	2	14 sequences to the Genus
<i>Recurvirostra avosetta</i>	4	2	2	0	8 sequences to the Genus
<i>Riparia riparia</i>	3	2	1	0	11 sequences to the Genus. 1 microsatellite for <i>R. cincta</i>
<i>Saxicola rubetra</i>	1	0	0	1	8 sequences to the Genus

Table 5. End.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Scolopax rusticola</i>	2	2	0	0	9 sequences to the Genus
<i>Sterna hirundo</i>	10	6	4	0	120 sequences to the Genus. 4 microsatellites for <i>S. dougallii</i>
<i>Streptopelia decaocto</i>	7	6	1	0	94 sequences to the Genus
<i>Streptopelia turtur</i>	7	6	1	0	idem
<i>Sturnus vulgaris</i>	28	14	14	0	32 sequences to the Genus
<i>Sylvia atricapilla</i>	12	6	6	0	54 sequences to the Genus
<i>Sylvia borin</i>	8	5	3	0	idem
<i>Sylvia cantillans</i>	3	3	0	0	idem
<i>Sylvia communis</i>	1	1	0	0	idem
<i>Sylvia conspicillata</i>	1	1	0	0	idem
<i>Sylvia hortensis</i>	1	1	0	0	idem
<i>Tachymarpis melba/</i> <i>Apus melba</i>	11	8	3	0	See <i>Apus</i>
<i>Tadorna tadorna</i>	9	7	2	0	22 sequences to the Genus
<i>Tichodroma muraria</i>	1	0	1	0	
<i>Tringa erythropus</i>	9	7	2	0	113 sequences to the Genus
<i>Tringa glareola</i>	11	9	2	0	idem
<i>Tringa nebularia</i>	10	8	2	0	idem
<i>Tringa ochropus</i>	11	8	3	0	idem
<i>Tringa totanus</i>	20	16	4	0	idem
<i>Turdus iliacus</i>	0	0	0	0	197 sequences to the Genus
<i>Turdus pilaris</i>	0	0	0	0	idem
<i>Turdus torquatus</i>	0	0	0	0	idem

ACKNOWLEDGMENTS

Thanks to L. M. Carrascal, I. Doadrio, M. García París and J. E. González for their commentaries and suggest. This work is partially supported by Comunidad de Madrid grant GR/AMB/0750/2004.

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Note in Proof:

Since the submission of this work and the proof corrections, and with respect to Aves, 4 microsatellites have been described for the genus *Porphyrio* and 16 for the species *Nycticorax nycticorax*. Many mitochondrial sequences have been published for the genus *Scolopax*. In the Reptilia, more sequences have been described for the genus *Blanus*, and 6 microsatellites have been described for *Mauremys*. Finally, many microsatellites have been described in the mammalian genus *Mustela*.

Table 6.— Sequences in database for mammals present in Comunidad de Madrid.

Tabla 6.— Secuencias en bases de datos para mamíferos presentes en la Comunidad de Madrid.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Apodemus sylvaticus</i>	372	yes	yes	21	1190 sequences to the Genus Several microsatellites
<i>Arvicola sapidus</i>	1	0	1	0	58 sequences to the Genus. 21 microsatellites for <i>A. terrestris</i>
<i>Barbastella barbastellus</i>	36	36	0	0	40 sequences to the genus
<i>Capra pyrenaica</i>	38	31	7	0	3740 sequences to the Genus. 55 microsatellites for <i>C. hircus</i>
<i>Capreolus capreolus</i>	358	yes	yes	16	383 sequences to the genus
<i>Cervus dama</i>	30	6	16	8	859 sequences to the Genus. 125 microsatellites
<i>Cervus elaphus</i>	405	yes	yes	82	idem
<i>Crociodura russula</i>	310	yes	yes	23	595 sequences to the Genus. 8 microsatellites for <i>C. suaveolens</i> . Genome mitochondrial completed
<i>Eliomys quercinus</i>	7	2	5	0	11 sequences to the Genus
<i>Eptesicus serotinus</i>	18	9	9	0	80 sequences to the Genus.
<i>Erinaceus europaeus</i>	185	96	77	12	9 microsatellites for <i>E. fuscus</i> 303 sequences to the Genus.
<i>Felis silvestris</i>	54	28	26	0	6313 sequences to the genus. 548 microsatellites
<i>Galemys pyrenaicus</i>	4	1	3	0	
<i>Genetta genetta</i>	27	22	5	0	125 sequences to the Genus
<i>Hypsugo savii</i>	18	18	0	0	20 sequences to the Genus
<i>Lepus granatensis</i>	104	51	53	0	1168 sequences to the Genus. 8 microsatellites for <i>L. saxatilis</i> and 6 for <i>L. europaeus</i>
<i>Lutra lutra</i>	118	81	10	27	130 sequences to the genus
<i>Lynx pardinus</i>	29	26	3	0	226 sequences to the Genus 6 microsatellites for <i>L. canadiensis</i> and 35 for <i>L. rufus</i> .
<i>Martes foina</i>	29	13	11	5	442 sequences to the Genus. 27 microsatellites
<i>Meles meles</i>	216	yes	yes	130	
<i>Microtus cabrerae</i>	34	2	32	0	1615 sequences to the Genus. 8 microsatellites for <i>M. oeconomus</i> and 8 for <i>M. montebelli</i>
<i>Microtus duodecimcostatus</i>	4	2	2	0	idem
<i>Microtus lusitanicus</i>	6	2	4	0	idem
<i>Microtus nivalis</i> / <i>Chionomys nivalis</i>	21	11	10	0	Idem (7 sequences to <i>Chionomys</i>)
<i>Miniopterus schreibersii</i>	123	110	8	5	171 sequences to the Genus
<i>Mus musculus</i>	8093572	yes	yes	more than 600	8099741 sequences to the Genus.
<i>Mustela nivalis</i>	55	48	5	2	585 sequences to the genus. 81 microsatellites
<i>Mustela putorius</i>	102	36	65	1	Idem
<i>Mustela vison</i>	206	yes	yes	63	idem
<i>Myotis bechsteini</i>	10	10	0	0	711 sequences to the Genus. 41 microsatellites for <i>M. myotis</i>
<i>Myotis blythii</i>	14	14	0	0	idem
<i>Myotis daubentoni</i>	42	17	25	0	idem
<i>Myotis emarginatus</i>	6	6	0	0	idem
<i>Myotis myotis</i>	168	30	97	41	idem
<i>Myotis mystacinus</i>	33	33	0	0	idem
<i>Myotis nattereri</i>	6	6	0	0	idem
<i>Neomys anomalus</i>	10	9	1	0	26 sequences to the Genus
<i>Nyctalus lasiopterus</i>	6	6	0	0	68 sequences to the Genus. 10 microsatellites to <i>N. noctula</i>
<i>Nyctalus leisleri</i>	8	8	0	0	idem
<i>Nyctalus noctula</i>	36	24	2	10	idem

Table 6. End.

Species	Sequences	Mitochondrial sequences	Nuclear sequences	Microsatellites	Remarks
<i>Oryctolagus cuniculus</i>	738692	yes	yes	More than 400	738795 sequences to the genus
<i>Ovis gmelini</i>	0	0	0	0	392342 sequences to the Genus. Many microsatellites
<i>Pipistrellus kuhli</i>	27	17	10	0	212 sequences to the Genus
<i>Pipistrellus nathusii</i>	13	13	0	0	idem
<i>Pipistrellus pipistrellus</i>	78	72	6	0	idem
<i>Plecotus auritus</i>	88	77	5	6	509 sequences to the Genus
<i>Plecotus austriacus</i>	54	54	0	0	Idem. Microsatellites for <i>P. auritus</i>
<i>Rhinolophus euryale</i>	1	1	0	0	1605 sequences to the Genus. Microsatellites for <i>R. ferrumequinum</i> and 14 for <i>R. hipposideros</i>
<i>Rhinolophus ferrumequinum</i>	1453	yes	yes	33	idem
<i>Rhinolophus hipposideros</i>	17	3	0	14	idem
<i>Rhinolophus mehelyi</i>	0	0	0	0	idem
<i>Sciurus vulgaris</i>	105	68	11	26	305 sequences to the Genus
<i>Sorex granarius</i>	2	2	0	0	662684 sequences to the Genus. 3 microsatellites for <i>S. cacutiens</i> , 3 for <i>S. unguiculatus</i> and 20 for <i>S. araneus</i>
<i>Sorex minutus</i>	75	75	0	0	idem
<i>Suncus etruscus</i>	2	2	0	0	62 sequences to the Genus
<i>Sus scrofa</i>	1113105	yes	yes	More than 2000	1113336 sequences to the Genus.
<i>Tadarida teniotis</i>	1	1	0	0	269 sequences to the Genus. 9 microsatellites for <i>T. brasiliensis</i>
<i>Talpa occidentalis</i>	2	1	1	0	45 sequences to the Genus
<i>Vulpes vulpes</i>	159	94	41	24	209 sequences to the Genus