

Linking Sequences with Specimens

Distichochlamys sp. AS18 chloroplast matK gene for maturase K, complete cds

GenBank: AB553309.1

[FASTA](#) [Graphics](#)Go to:

LOCUS AB553309 1542 bp DNA linear PLN 02-OCT-2010

DEFINITION Distichochlamys sp. AS18 chloroplast matK gene for maturase K, complete cds.

ACCESSION AB553309

VERSION AB553309.1 GI:308072161

KEYWORDS .

SOURCE chloroplast Distichochlamys sp. AS18

ORGANISM [Distichochlamys sp. AS18](#)

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Zingiberales; Zingiberaceae; Distichochlamys.

REFERENCE 1

AUTHORS Takano,A. and Nagamasu,H.

TITLE Monophyly of Myxochlamys (Zingiberaceae) is confirmed: Molecular phylogenetic analysis

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 1542)

AUTHORS Takano,A. and Nagamasu,H.

TITLE Direct Submission

JOURNAL Submitted (01-APR-2010) Contact:Atsuko Takano Museum of Nature and Human Activities, Hyogo; 6 chome, Yayoigaoka, Sanda, Hyogo 669-1546, Japan

FEATURES Location/Qualifiers

source

1..1542

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Change region shown Customize view Analyze this sequence [Run BLAST](#)[Pick Primers](#)[Highlight Sequence Features](#)[Find in this Sequence](#)LinkOut to external resources [RBGE Herbarium Catalogue E00198079](#)
[Royal Botanic Garden Edinburgh][RBGE Living Collection 20021433](#)
[Royal Botanic Garden Edinburgh]Related information [Related Sequences](#)[Protein](#)[Taxonomy](#)Recent activity [Turn Off](#) [Clear](#) [Distichochlamys sp. AS18 chloroplast matK gene for m](#); Nucleotide [The absence of sharks from abyssal regions of the world's oceans](#) PMC [txid473364\[Organism:noexp\] \(1\)](#) PMC [Etmostegus princeps](#)



Accession: 20021433 - *Distichoclamys* by: Helen A. Yeats on 23/2/2011
Image 1 of 2

CLOSE X

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Lesula: a new species of *Cercopithecus* monkey endemic to the Democratic Republic of Congo and implications for conservation of Congo's central basin.

Hart JA, Detwiler KM, Gilbert CC, Burrell AS, Fuller JL, Emetsu M, Hart TB, Vosper A, Sargis EJ, Tosi AJ.

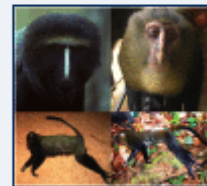
Lukuru Wildlife Research Foundation, Kinshasa, Gombe, Democratic Republic of Congo ; Division of Vertebrate Zoology, Yale Peabody Museum of Natural History, New Haven, Connecticut, United States of America.

Abstract

In June 2007, a previously undescribed monkey known locally as "lesula" was found in the forests of the middle Lomami Basin in central Democratic Republic of Congo (DRC). We describe this new species as *Cercopithecus lomamiensis* sp. nov., and provide data on its distribution, morphology, genetics, ecology and behavior. *C. lomamiensis* is restricted to the lowland rain forests of central DRC between the middle Lomami and the upper Tshuapa Rivers. Morphological and molecular data confirm that *C. lomamiensis* is distinct from its nearest congener, *C. hamlyni*, from which it is separated geographically by both the Congo (Lualaba) and the Lomami Rivers. *C. lomamiensis*, like *C. hamlyni*, is semi-terrestrial with a diet containing terrestrial herbaceous vegetation. The discovery of *C. lomamiensis* highlights the biogeographic significance and importance for conservation of central Congo's interfluvial TL2 region, defined from the upper Tshuapa River through the Lomami Basin to the Congo (Lualaba) River. The TL2 region has been found to contain a high diversity of anthropoid primates including three forms, in addition to *C. lomamiensis*, that are endemic to the area. We recommend the common name, lesula, for this new species, as it is the vernacular name used over most of its known range.

PMID: 22984482 [PubMed - indexed for MEDLINE] PMCID: PMC3440422 [Free PMC Article](#)

Images from this publication. [See all images \(10\)](#) [Free text](#)



[+](#) **Publication Types, MeSH Terms**

[+](#) **LinkOut - more resources**

Cercopithecus lomamiensis

Taxonomy ID: 1191211

Genbank common name: **lesula**

Inherited blast name: **primates**

Rank: species

Genetic code: [Translation table 1 \(Standard\)](#)

Mitochondrial genetic code: [Translation table 2 \(Vertebrate Mitochondrial\)](#)

Other names:

synonym: **Cercopithecus sp. ASB-2012**

authority: **Cercopithecus lomamiensis Hart et al. 2012**

type material: **YPM MAM 14189**

type material: **YPM MAM 14191**

type material: **YPM MAM 14192**

type material: **YPM MAM 14080**

type material: **YPM 14191**

type material: **YPM 14080**

type material: **YPM 14192**

type material: **YPM 14189**

Lineage(full)

[cellular organisms](#); [Eukaryota](#); [Opisthokonta](#); [Metazoa](#); [Eumetazoa](#); [Bilateria](#); [Deuterostomia](#); [Chordata](#); [Craniata](#); [Vertebrata](#); [Gnathostomata](#); [Teleostomi](#); [Euteleostomi](#); [Sarcopterygii](#); [Tetrapoda](#); [Amniota](#); [Mammalia](#); [Theria](#); [Eutheria](#); [Euarchontoglires](#); [Primates](#); [Haplorrhini](#); [Simiiformes](#); [Catarrhini](#); [Cercopithecoidea](#); [Cercopithecidae](#); [Cercopithecinae](#); [Cercopithecus](#)

Comments and References:



[Hart et al. \(2012\)](#)

External Information Resources (NCBI LinkOut)

LinkOut	Subject	LinkOut Provider
search Catalog of Life	taxonomy/phylogenetic	Catalog of Life
search GBIF	taxonomy/phylogenetic	Global Biodiversity Information Facility
Cercopithecus (guenon)	taxonomy/phylogenetic	Primate Fact Sheets

Entrez records

Database name	Direct links
Nucleotide	8
Protein	4
Taxonomy	1

Display Settings: Summary, Sorted by Default order

Results: 4

[Cercopithecus lomamiensis isolate ME408 X chromosome intergenic region genomic sequence](#)

1. 4,688 bp linear DNA

Accession: JN106060.1 GI: 387865320

[GenBank](#) [FASTA](#) [Graphics](#) [Related Sequences](#) [Full text in PMC](#) [PubMed](#) [Taxonomy](#)

[Cercopithecus lomamiensis isolate GP609 X chromosome intergenic region genomic sequence](#)

2. 4,819 bp linear DNA

Accession: JN106063.1 GI: 387865323

[GenBank](#) [FASTA](#) [Graphics](#) [Related Sequences](#) [Full text in PMC](#) [PubMed](#) [Taxonomy](#)

[Cercopithecus lomamiensis isolate ME408 testis-specific protein \(TSPY\) gene, partial cds](#)

3. 2,237 bp linear DNA

Accession: JN106057.1 GI: 387865316

[GenBank](#) [FASTA](#) [Graphics](#) [Related Sequences](#) [Full text in PMC](#) [Protein](#) [PubMed](#) [Taxonomy](#)

[Cercopithecus lomamiensis isolate GP609 testis-specific protein \(TSPY\) gene, partial cds](#)

4. 2,237 bp linear DNA

Accession: JN106055.1 GI: 387865312

[GenBank](#) [FASTA](#) [Graphics](#) [Related Sequences](#) [Full text in PMC](#) [Protein](#) [PubMed](#) [Taxonomy](#)

Display Settings: Summary, Sorted by Default order

Cercopithecus lomamiensis isolate ME408 X chromosome intergenic region genomic sequence

GenBank: JN106060.1

[FASTA](#) [Graphics](#)

[Go to:](#)

LOCUS JN106060 4688 bp DNA linear PRI 05-MAR-2013
DEFINITION Cercopithecus lomamiensis isolate ME408 X chromosome intergenic region genomic sequence.
ACCESSION JN106060
VERSION JN106060.1 GI:387865320
KEYWORDS .
SOURCE Cercopithecus lomamiensis (lesula)
ORGANISM [Cercopithecus lomamiensis](#)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Haplorrhini; Catarrhini; Cercopithecidae; Cercopithecinae; Cercopithecus.
REFERENCE 1 (bases 1 to 4688)
AUTHORS Hart,J.A., Detwiler,K.M., Gilbert,C.C., Burrell,A.S., Fuller,J.L., Emetshu,M., Hart,T.B., Vosper,A., Sargis,E.J. and Tosi,A.J.
TITLE Lesula: A New Species of Cercopithecus Monkey Endemic to the Democratic Republic of Congo and Implications for Conservation of Congo's Central Basin
JOURNAL PLoS ONE 7 (9), E44271 (2012)
PUBMED [22984482](#)
REMARK Publication Status: Online-Only
REFERENCE 2 (bases 1 to 4688)
AUTHORS Hart,J.A., Detwiler,K.M., Gilbert,C.C., Burrell,A.S., Fuller,J.L., Emetshu,M., Hart,T.B., Vosper,A., Sargis,E.J. and Tosi,A.J.
TITLE Direct Submission
JOURNAL Submitted (10-JUN-2011) Department of Anthropology, New York University, 25 Waverly Place, New York, NY 10003, USA
FEATURES Location/Qualifiers
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YPM Mammalogy - Online Catalog



2 Apr 2013 15:39:29

Items 1-1 of 1 matching items.


[New Search](#)

YPM MAM 014080	
Taxon Name..... <i>Cercopithecus lomamiensis</i> J. Hart, Detwiler, Gilbert, Burrell, Fuller, Emetshu, T. Hart, Vosper, Sargis, and Tosi, 2012 - HOLOTYPE	
Locality..... Africa. Democratic Republic of Congo. Orientale Province. Tshopo District. Lohumonoko. shot. Elev. 470 m.	
LatLon..... -1.02237 24.42368	n=1
Collected..... M. Emetshu. 12 Aug 2008.	
Higher Ranks..... Primates; Cercopithecidae	
Common Name..... Lesula	
Other Attributes... TISSUE; SKIN; male male; adult adult; skin, flat (tissue sample at at New York University); skeleton (skull only); TYPE; SKELETON	

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Comments to: peabody.webmaster@yale.edu

Institution ID	281
Full institution name	Museum of Comparative Zoology, Harvard University
Institution code	MCZ
Also known as	
Country	USA
Address	26 Oxford Street Cambridge, Massachusetts 02138
Institution phone	617-495-3045
Institution fax	
Record source	
Institution web page	http://www.mcz.harvard.edu/
Url rule	
Comments	= MCZC
Collection type	museum
Qualifier type	specimen_voucher 
Collection code	HERP A R I
Contact person name	Linda S. Ford
Contact person e-mail	lford@oeb.harvard.edu
Contact person phone	(617) 495-3748
Contact person fax	(617) 495-5667
Contact comments	

Institution ID	281
Full institution name	Museum of Comparative Zoology, Harvard University
Institution code	MCZ
Also known as	
Country	USA
Address	26 Oxford Street Cambridge, Massachusetts 02138
Institution phone	617-495-3045
Institution fax	
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Etmopterus princeps voucher KU:IT:T3681 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial

GenBank: FJ519565.1

[FASTA](#) [Graphics](#)

Go to:

LOCUS FJ519565 652 bp DNA linear VRT 03-APR-2013

DEFINITION Etmopterus princeps voucher KU:IT:3681 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial.

ACCESSION FJ519565

VERSION FJ519565.1 GI:261875099

KEYWORDS BARCODE.

SOURCE mitochondrion Etmopterus princeps (great lanternshark)

ORGANISM [Etmopterus princeps](#)

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes; Elasmobranchii; Squalimorphii; Squaliformes; Etmopteridae; Etmopterus.

REFERENCE 1 (bases 1 to 652)

AUTHORS Wong,E.H., Shivji,M.S. and Hanner,R.H.

TITLE Identifying sharks with DNA barcodes: assessing the utility of a nucleotide diagnostic approach

JOURNAL Mol Ecol Resour 9 (SUPPL S1), 243-256 (2009)

PUBMED [21564984](#)

REFERENCE 2 (bases 1 to 652)

AUTHORS Wong,E.H.-K.

TITLE Direct Submission

JOURNAL Submitted (04-DEC-2008) Integrative Biology, University of Guelph, 50 Stone Rd E, Guelph, Ontario N1G2W1, Canada

FEATURES Location/Qualifiers

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Hypoplectrus floridae voucher hecosur1 cytochrome oxidase subunit 1 (COI) gene, partial cds; mitochondrial

GenBank: JX444756.1

[FASTA](#) [Graphics](#)

[Go to:](#)

LOCUS JX444756 652 bp DNA linear VRT 03-APR-2013
DEFINITION Hypoplectrus floridae voucher SIO:12-58 cytochrome oxidase subunit 1 (COI) gene, partial cds; mitochondrial.
ACCESSION JX444756
VERSION JX444756.1 GI:403488715
KEYWORDS BARCODE.
SOURCE mitochondrion Hypoplectrus floridae
ORGANISM [Hypoplectrus floridae](#)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Percoidei; Serranidae; Serraninae; Hypoplectrus.
REFERENCE 1 (bases 1 to 652)
AUTHORS Victor,B.C.
TITLE Hypoplectrus floridae n. sp. and Hypoplectrus ecosur n. sp., two new Barred Hamlets from the Gulf of Mexico (Pisces: Serranidae): more than 3% different in COI mtDNA sequence from the Caribbean Hypoplectrus species flock
JOURNAL J Ocean Sci Found 5, 1-19 (2012)
REFERENCE 2 (bases 1 to 652)
AUTHORS Victor,B.
TITLE Direct Submission
JOURNAL Submitted (01-AUG-2012) Biology, Ocean Science Foundation, 4051 Glenwood St., Irvine, CA 92604, USA
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Hypoplectrus floridae, n. sp.

Figs. 1–5

Holotype. SIO 12-58 (1) 40.3 mm SL, SW Florida, off Everglades, Ten Thousand Islands (25.6°, -81.25°), July 14, 2011, SEAMAP surveys.

Paratypes. SIO 12-58 (3) 33.4–39.4 mm SL, same as holotype; SIO 12-59 (1) 33.7 mm SL, SW Florida, off Everglades, Ten Thousand Islands (25.7°, -81.34°), July 13, 2011, SEAMAP surveys.

Diagnosis. A hamlet with the usual color and patterns of the brown Barred Hamlet but with several distinguishing markings: a pair of dark rounded spots at the base of the caudal fin placed above and below the midline, usually symmetrical, present on all stages, including large adults; a break in the narrow mid-body bar (the fourth bar, after the wide mid-body bar) just above the lateral line, usually associated with well-delineated and unbroken bars to each side; frequently a short rearward spur at the top of the last body bar (the fifth bar, forward of the caudal-peduncle bar), usually outlining a light wedge just below the base of the last dorsal-fin rays. Characters not frequent on Caribbean Barred Hamlets, but typically associated with *H. floridae* (and *H. ecosur*) include the bar under the eye being orange, even when the body bars are brown, and a dusky pelvic fin.

Description. Body wide and broadly oval, maximum body depth just behind operculum 38–42 (42)% SL (range of paratypes (holotype)), and compressed, side-to-side width 12–15 (13)% SL (measured just forward of pectoral-fin base); predorsal length 38–42 (42)% SL; preanal length 65–71 (67)% SL; prepelvic length 36–41 (40)% SL; caudal peduncle depth 14–15 (15)% SL, caudal peduncle length (dorsal) 7–10 (8)% SL; lateral line complete, curving in a high arch over pectoral fin becoming straight on caudal peduncle.

Head large 41–43 (41)% SL; dorsal head profile smooth and mostly straight, rising sharply from terminal tip of jaw to dorsal-fin origin; maximum head depth (measured at the rear end of the operculum) 37–41 (42)% SL; eyes large and round, orbit diameter 26–29 (28)% HL, pupil pear-shaped, pointing forward; interorbital space flat and relatively narrow, minimum width 13–15 (14)% HL; snout sharply pointed and short 23–30 (29)% HL; upper preopercular margin tilted slightly forward of vertical with a rounded angle to lower limb, small regular straight serrations on both limbs, longest at angle, about 22–33 in total; opercle with three flat spines, the middle largest and at the level of the lower third of eye. Anterior nostril a short tube, posterior nostril a flat opening with a diameter about half intermarial distance.



Figure 1. *Hypoplectrus floridae*, holotype, SIO 12-58, 40.3 mm SL, Florida, Ten Thousand Islands.

Hypoplectrus floridae

Taxonomy ID: 1230408

Inherited blast name: **bony fishes**

Rank: species

Genetic code: [Translation table 1 \(Standard\)](#)

Mitochondrial genetic code: [Translation table 2 \(Vertebrate Mitochondrial\)](#)

Other names:

authority: **Hypoplectrus floridae Victor, 2012**

type material: **SIO 12-58 (lot)**

type material: **hecosur1**

equivalent name: **Hypoplectrus sp. BV-2012**

Lineage(abbreviated)

[Eukaryota](#); [Metazoa](#); [Chordata](#); [Craniata](#); [Vertebrata](#); [Euteleostomi](#);
[Actinopterygii](#); [Neopterygii](#); [Teleostei](#); [Euteleostei](#); [Neoteleostei](#);
[Acanthomorpha](#); [Acanthopterygii](#); [Percomorpha](#); [Perciformes](#); [Percoidei](#);
[Serranidae](#); [Serraninae](#); [Hypoplectrus](#)

Comments and References:

Victor (unpublished, 2012)

Victor,B. "A new hamlet from the Gulf of Mexico (Perciformes:Serranidae:Hypoplectrus)" Unpublished (as of 6 Sep 2012)

 [Victor \(2012\)](#)

Victor,B.C. (2012) "Hypoplectrus floridae n. sp. and Hypoplectrus ecosur n. sp., two new Barred Hamlets from the Gulf of Mexico (Pisces: Serranidae): more than 3% different in COI mtDNA sequence from the Caribbean Hypoplectrus species flock" Journal of the Ocean Science Foundation 5: 1-19

External Information Resources (NCBI LinkOut)

LinkOut	Subject	LinkOut Provider
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search GBIF	taxonomy/phylogenetic	Global Biodiversity Information Facility

Entrez records

Database name	Direct links
Nucleotide	5
Protein	5
Taxonomy	1