

# DNA Barcodes of Moths (Lepidoptera) from Lake Turkana, Kenya

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#### Note

## DNA barcodes of moths (Lepidoptera) from Lake Turkana, Kenya

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This paper provides metadata for DNA barcode (COI) data in GenBank for a collection of moths (Lepidoptera) made at South Turkwel near Lake Turkana, Kenya. This paper aims to make DNA barcode data available to document ongoing research, to contribute to the International Barcode of Life (iBOL; www. ibol.org) and Kenya Barcode of Life projects, and to encourage enhancement in identifications, in line with the concept of DNA barcode data release papers and the Fort Lauderdale principles for genetic data (Schindel et al. 2011). Data for 89 sequences representing 35 barcode clusters (putative species) have been released on GenBank (accession numbers KF147250-KF147332, KF603887) including the required fields for the BARCODE data standard (Benson et al. 2012) and more data, including images, are available on BOLD (www.boldsystems. org; Ratnasingham and Hebert 2007), accessible from the project TBILE using a DOI (dx.doi.org/10.5883/DS-TBILE).

There is very little literature on the moths of the Turkana Basin. Probably the best known collection was made by the 1934 Lake Rudolf Rift Valley Expedition, but the results were never published, although we have seen specimens in Natural History Museum, London (BMNH) (Buxton 1936, Vári 1964). We have also seen Turkana moth specimens in the National Museums of Kenya (NMK), but there is no comprehensive published documentation. We have been able to identify some of the moths by comparison to collections at the Smithsonian National Museum of Natural History (USNM),

BMNH, and NMK, the literature, or matching DNA sequences in BOLD. However, because of the poor state of knowledge of African Lepidoptera (e.g., Mey 2011), further refinement will take considerable time. Thus, we are making these data available now, while we continue the process of identifications. Where taxonomic names are not readily available from existing literature, the DNA cluster-based morphospecies can be used as species hypotheses that can confirmed by future taxonomic studies in broader context of the African fauna (Schindel and Miller 2010, Ratnasingham and Hebert 2013).

## MATERIALS AND METHODS

Sixty-six of the specimens were collected at the Turkana Basin Initiative (TBI) station at West Turkana, 465 m, on the Turkwel River, 29 km east of Lodwar, near Lake Turkana, 3.144° N, 35.863° E, 14-18 August 2011. Most specimens were collected at lights, and all were killed by freezing and dried as pinned voucher specimens. Four specimens of Eoophyla excentrica Mey & Speidel (Crambidae: Acentropinae) were collected offshore Central Island, Lake Turkana, 360 m, 3.507° N, 36.033° E, 16 August 2011. Comparative specimens are included from Kenya (6), Nigeria (10), Papua New Guinea (2), and the United States (1).

Genitalic dissections follow Robinson (1976). Morphological comparisons were made to both the literature and to collections of USNM, BMNH, and NMK. Additional context was provided by intensive sampling of moths from light at Mpala Research Centre in central Kenya,

from 1998 to 2011 by Scott Miller and Tina Kuklenski (Adamski et al. 2010). Vouchers are retained by USNM and NMK. DNA sequencing (COI barcode) followed standard methods at the Biodiversity Institute of Ontario, University of Guelph, between 2008 and 2012 (Craft et al. 2010, Hrcek et al. 2011, Wilson 2012), using legs from pinned moths. Up to five specimens per morphospecies were sampled (9 in one case because of marked variation in wing pattern). All of the Turkana vouchers sampled for DNA yielded successful sequences, all but one of which yielded a full length barcode meeting the BARCODE keyword standard in GenBank (Benson et al. 2012). Full details on the sequences are provided in BOLD (Ratnasingham and Hebert 2013) through DOI (dx.doi.org/ 10.5883/DS-TBILE). Barcode clusters are based on the RESL algorithm as implemented in BOLD as described by Ratnasingham and Hebert (2013).

The climate of this region is arid and considered semi-desert with > 250 mm of rain falling annually. Temperatures are high and maximum daily highs typically exceed 40 °C and occasionally 50 °C. The vegetation at the collection sites is dominated by the trees *Acacia tortilis* and *Acacia reficiens*, shrubs *Salvadora persica* and *Cadaba rotundifolia*, and perennial herbaceous *Indigofera spinosa*, with annual *Aristida* spp. grasses occurring seasonally after rains.

## RESULTS

Identifications for all specimens are provided on BOLD (dx.doi.org/10.5883/DS-TBILE) and GenBank. While many specimens remain unidentified, some identifications are worth comment here:

## Cosmopterigidae: Chrysopeleiinae

Bifascioides leucomelanella (Rebel) (male genitalia slide USNM 125903);

previously recorded from Kenya based only on a specimen collected in 1935 inside an Imperial Airways airplane at Kisumu (Kasy 1968: 513).

## Crambidae: Spilomelinae

Marasmia trapezalis Guenée; DNA barcode results confirm its status as a cosmopolitan species (Clarke 1971) as specimens from Nigeria (GenBank KF147300, 147304-147312), Papua New Guinea (GenBank KF147302-KF147303) and Florida (GenBank KF603887) possess almost identical DNA sequences. Barcode cluster AAC0297 in BOLD also includes specimens from Mexico, Australia, Sierra Leone, South Africa (BOLD, unpublished data), and Palau (GenBank JX017849, Haines and Rubinoff 2012). This species is the major rice leaf folder in West Africa (Heinrichs and Barrion 2004: 63), and is a pest of various graminaceous crops in Asia (Mathew and Menon 1986, Khan et al. 1988). This species is often placed in the genus Cnaphalocrocis, but we follow Munroe (1991) in placing it in Marasmia. Buettiker and Gallagher (1980) comment on the biology of the species in Oman.

### Erebidae: Erebinae

Gnamptonyx innexa (Walker); widespread from North Africa to India, but not previously recorded from Kenya (Hacker et al. 2010: 22).

While the sample size is too small for serious biogeographic analysis, some interesting trends are evident from comparison to other barcode clusters in BOLD, including an extensive sampling of about 2000 moths from Mpala, Kenya, representing over 700 barcode clusters. Of the 34 species from the TBI station near Lodwar, none are represented in BOLD from Mpala, or from Serengeti, Tanzania (Hebert, unpublished data in BOLD). However, two of the species recorded at

TBI are also known from Nguruman in southern Kenya (1.848° S, 36.1° E, 660 m), including *Dysodia lutescens* Whalley 1968 (Thyrididae), and an unidentified Phycitinae. Interestingly, five also occur in the United Arab Emirates (E. van Nieukerken, unpublished data in BOLD) and three others occur in Yemen (A. Hausmann, unpublished data in BOLD). Thus, the fauna at Turkana has a distinctly North African relationship.

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