The 'cyber-centipede':

The first eukaryotic organism described with a fully sequenced transcriptome, DNA barcode and micro computed tomography



L. Penev, P. Stoev, A. Komerički, N. Akkari, S. Li, X. Zhou, S. Edmunds, C. Hunter, A.M. Weigand, D. Porco, M. Zapparoli, T. Georgiev, D. Mietchen, D. Roberts, V. Smith

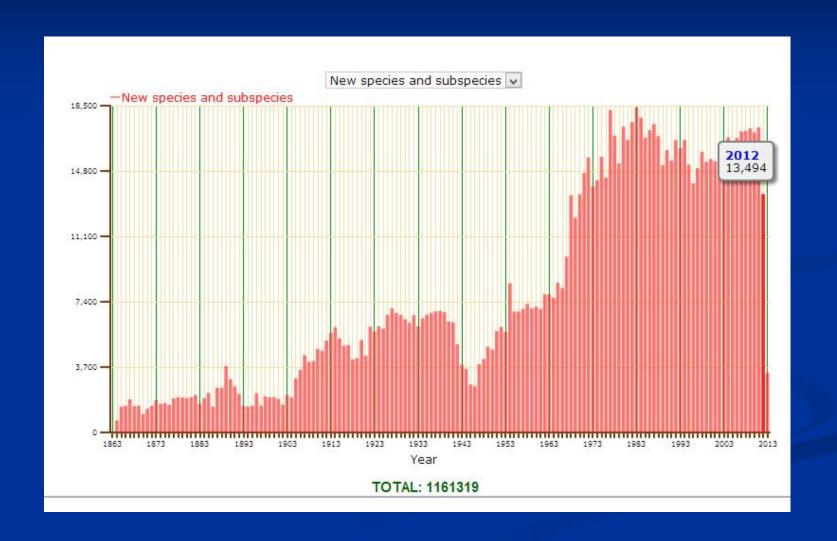




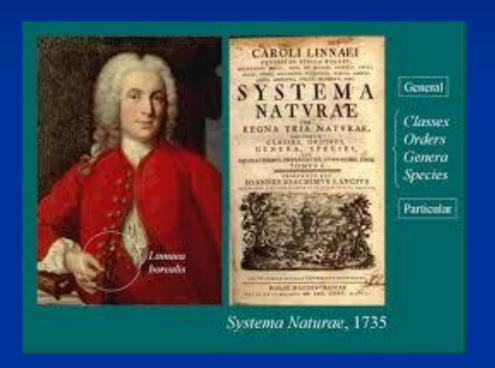




"Only" 13,494 new animal species described in 2012 (Index of Organism Names)

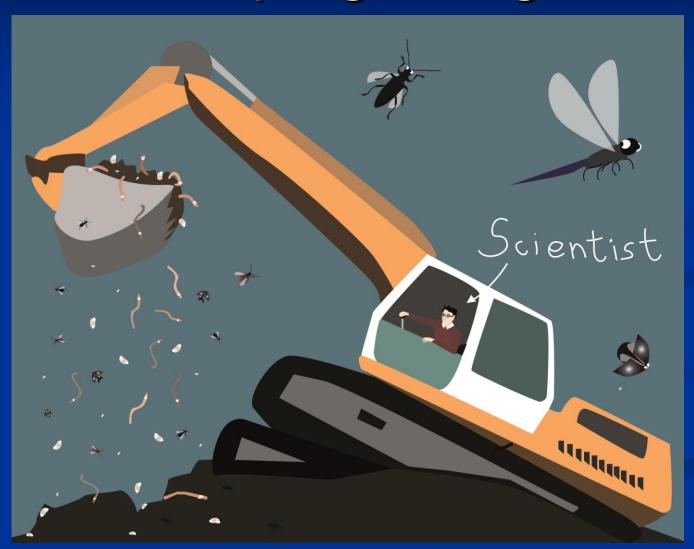


How much have taxonomic descriptions changed since Linnaeus?



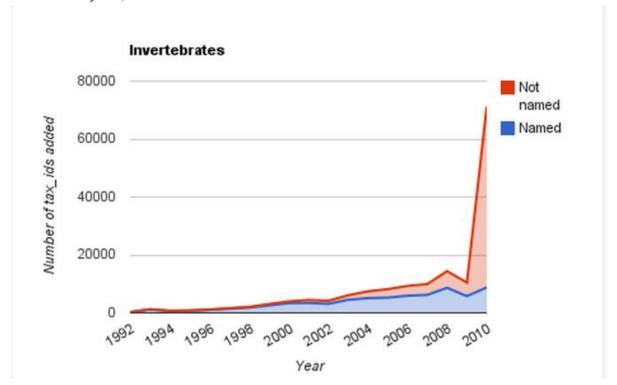
273. SCOLOPENDRA. Antennae setaceae. Palpi duo filiformes, articulati, inter maxillas con-Labium dentatum fissum. Corpus depressum. Pedes numerosi, totidem utrinque, quot corporis segmenta. I. Sc. pedibus utrinque duodecim, corpore ovali, cauda penicillo albo. Fn. fuec. 2063.* Geoffr. inf. par. 2. p.677. n. 6. t. 22. f. 4. Fabr. fp. inf. 1. p. 531. n. 1. mant. inf. I. p. 341. n. I. Scop. ent. carn. 1152. Julus lagurus. Degeer inf. 7. p. 571. n. 1. t. 36. f. 1 - 3. Julus penicillatus. AEt. par. mill. 1. p. 532. t. 17. f. 4-10. Habitat fub Europae mufcis.

The data deluge Massive sampling/data gathering



Proportion of dark taxa increase dramatically!

For "invertebrates" 2010 saw an explosive growth in the number of new taxa sequenced, with nearly 71,000 new taxa added to GenBank.



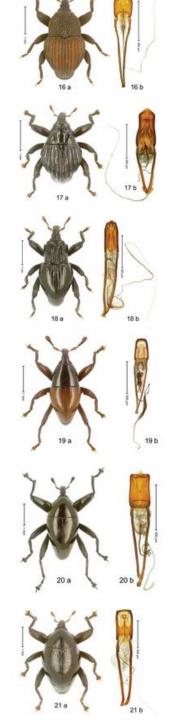
This coincides with a spectacular drop in the number of properly-named taxa, but even before 2010 the proportion of named invertebrate species in GenBank was in decline: in 2009 just over a half of the species added had binomials.

Rod Page, iPhylo blogspot, 12 April 2011

Low uptake of open data publishing/sharing



Solutions?



Turbo-taxonomy: publish fast and many!

Pipeline descriptions of new species (Butcher et al. 2012, Riedel et al. 2013b):

- molecular data
- concise morphological descriptions
- digital imaging

66. *Trigonopterus ptolycoides* Riedel, sp. n. urn:lsid:zoobank.org:act::37E45438-EBB4-4FD0-BC0E-B72E4D9979F8 http://species-id.pet/wiki/Trigonopterus ptolycoides

Diagnostic description. Holotype, male (Fig. 66a). Length 4.28 mm. Color ferruginous, elytral disk with some irregular darker spots; integument partly covered with yellowish-brownish tomentum. Body dorsally flattened, with irregular lateral ridge; with constrictions at middle of pronotum, between pronotum and elytron, and in basal third of elytron. Rostrum in basal half with distinct median ridge and pair of submedian ridges; basally above eyes with pair of protrusions; apical ½ subglabrous, punctulate. Pronotum with subapical constriction; with marked lateral flanges; coarsely rugose-punctate; punctures containing each one upcurved scale and much finer tomentum. Elytra with distinct striae; intervals with irregular tomentose tubercles; with marked lateral flanges; base bisinuate; apex extended ventrad, slightly beak-shaped. Femora narrow, parallel-sided, edentate. Meso- and metafemur dorsally with fringe of erect scales; metafemur subapically with stridulatory patch. Meso- and metatibia tapering from base to apex. Tarsi asymmetrical; tarsomere 3 with anterior lobe much larger than posterior lobe. Abdominal ventrite 5 apically deeply emarginate. Aedeagus (Fig. 66b) with body flattened, sides subparallel, apex pointed; transfer-apparatus short, spiniform; ductus ejaculatorius without bulbus. Intraspecific variation. Length 3.92–4.28 mm. Female abdominal ventrite 5 apically rounded.

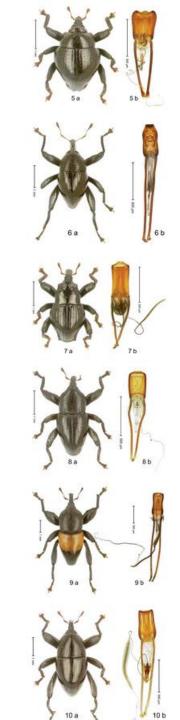
Material examined. Holotype (SMNK): ARC1415 (EMBL # HE615964), PAPUA NEW GUINEA, Eastern Highlands Prov., 37 km S Goroka, Hogave vill., Mt. Michael, S06°22.798', E145°15.427' to S06°22.925', E145°16.645', 2179–2800 m, 09-15-VII-2009, sifted. Paratypes (NAIC, SMNK, ZSM): 6 exx, ARC1416 (EMBL # HE615965), ARC1417 (EMBL # HE615966), same data as holotype.

Distribution. Eastern Highlands Prov. (Mt. Michael). Elevation: ca. 2179–2800 m.

Biology. Sifted from leaf litter in primary forest.

Etymology. This epithet is a combination of the genus name *Ptolycus* and the Latin suffix *-oides* (having the form of) and refers to the species' resemblance in habitus.

Notes. Trigonopterus ptolycoides Riedel, sp. n. was coded as "Trigonopterus sp. 68" by Tänzler et al. (2012).







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Biodiversity Data Journal 1: e1013 (28 Oct 2013) doi: 10.3897/BDJ.1.e1013







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Taxonomic paper

Eupolybothrus cavernicolus Komerički & Stoev sp. n. (Chilopoda: Lithobiomorpha: Lithobiidae): the first eukaryotic species description combining transcriptomic, DNA barcoding and micro-CT imaging data

▼ Pavel Stoev, Ana Komerički, Nesrine Akkari, Shanlin Liu, Xin Zhou, Alexander M. Weigand, Jeroen Hostens, Christopher I. Hunter, Scott C. Edmunds, David Porco, Marzio Zapparoli, Teodor Georgiev, Daniel Mietchen, David Roberts, Sarah Faulwetter, Vincent Smith, Lyubomir Penev

Abstract

We demonstrate how a classical taxonomic description of a new species can be enhanced by applying new generation molecular methods, and novel computing and imaging technologies. A cave-dwelling centipede, Eupolybothrus cavernicolus Komerički & Stoev sp. n. (Chilopoda: Lithobiomorpha: Lithobiidae), found in a remote karst region in Knin, Croatia, is the first eukaryotic species for which, in addition to the traditional morphological description, we provide a fully sequenced transcriptome, a DNA barcode, detailed anatomical X-ray microtomography (micro-CT) scans, and a movie of the living specimen to document important traits of its ex-situ behaviour. By employing micro-CT scanning in a new species for the first time, we create a high-resolution morphological and anatomical dataset that allows virtual reconstructions of the specimen and subsequent interactive manipulation to test the recently introduced 'cybertype' notion. In addition, the transcriptome was recorded with a total of 67,785 scaffolds, having an average length of 812 bp and N50 of 1,448 bp (see GigaDB). Subsequent annotation of 22,866 scaffolds was conducted by tracing homologs against current available databases, including Nr, SwissProt and COG. This pilot project illustrates a workflow of producing, storing, publishing and disseminating large data sets associated with a description of a new taxon. All data have been deposited in publicly accessible repositories, such as GigaScience GigaDB, NCBI, BOLD, Morphbank

Article title Abstract

Introduction

The study group

Materials and methods

Collected material and morphological study Molecular experiments and sequencing

Micro-CT scanning Abbreviations

Taxon treatments

Eupolybothrus cavernicolus

Eupolybothrus leostygis

Identification key to the species of Eupolybothrus (Schizopolybothrus) based on adult males

Analysis

Molecular delimitations

Transcriptome analysis and annotation

Discussion

Taxonomic affinities

Micro-computed tomography and 'cybertype' notion

Data management and release

Conclusions Acknowledgements

References

Eupolybothrus cavernicolus



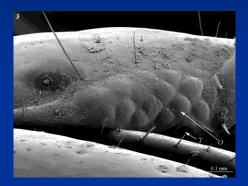
Methods

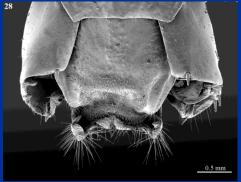
- Morphological study with a Zeiss microscope
- SEM images
- Mitochondrial
 Cytochrome C Oxidase
 Subunit I gene
 sequencing
- Full transcriptome sequencing
- Micro-CT scanning
- Video of living specimen



Morphology

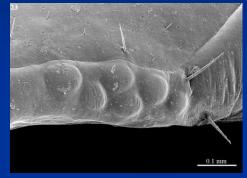
E. cavernicolus sp.n.



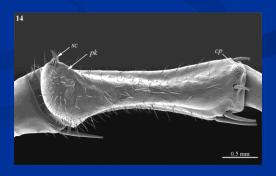




E. leostygis

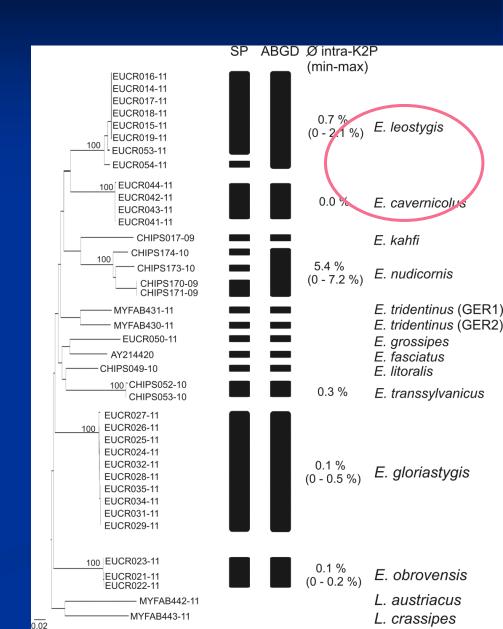






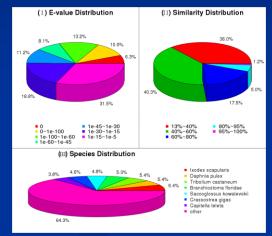
Molecular delimitation

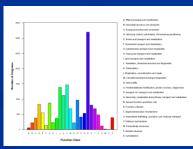
- 1. The final dataset comprises 39 COI sequences from 10 species.
- Molecular delimitation through ABGD algorithm - Puillandre et al. (2012)
- 3. The new species clusters with *E. leostygis*, its morphologically closest neigbour
- 4. Intraspecific K2P genetic distance was 0.0 % for *E. cavernicolus*

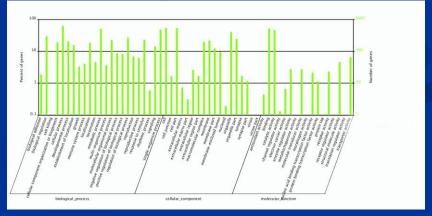


Transcriptome sequencing

- i. A total of 67,785 scaffolds were produced with an average length of 812 bp.
- ii. Subsequent annotation was conducted by tracing homologs against current available databases
- iii. Using orthoDB, 2,188 one to one orthologs were filtered out from four selected arthropod genomes: Drosophila melanogaster, Daphnia pulex, Ixodes scapularis and Strigamia maritima







Micro-CT in Taxonomy

By employing micro-CT scans in taxonomy, important morphological and anatomical characters can be examined in their natural position without damage to the original specimen

Micro-CT scan

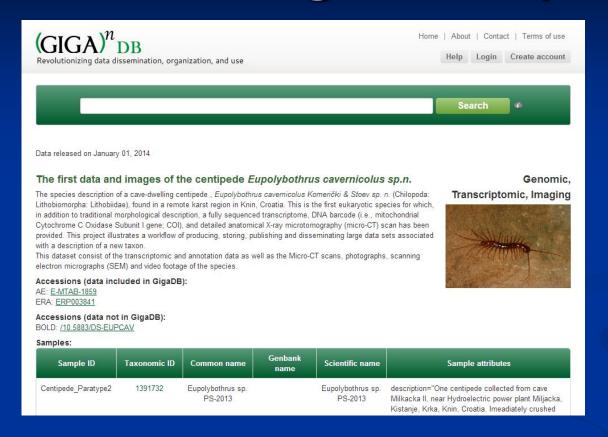


Faulwetter et al. (2013) suggested the creation of high-resolution virtual morphological and anatomical data libraries allowing reconstruction and interactive manipulation of type specimens, the so-called 'cybertypes'.

Data repositories

- GigaDB (complete database)
- NCBI (COI and transcriptome sequences)
- BOLD (COI)
- Morphbank (morphological images)
- Morphosource (micro CT row data, 3D model)
- Plazi (taxon treatments of legacy literature related to genus *Eupolybothrus*

Data management pilot



GigaDB in this example provides direct links between the genomic and micro-CT data, through a Darwin Core CSV dataset describing the type specimens, as well as capturing all of the metadata in the interoperable ISA-TAB format

Next generation sequencing and imaging require next generation publishing!

- Focus and Scope
- Globally unique innovations
- · Criteria for publication.
- Peer review
- For authors
- Data publication
- Publication fees . Licenses and Copyright
- Frequently Asked Questions (FAQ)
- Contacts

Editorial team











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authoring

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Data and text publishing

integrated

<u>Dr Lyubomir Penev</u> ▼

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Increase collaboration

Respond to funding requirements



Establish scientific priority







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- Large datasets published as data papers
- Text stored and treated as data
- Community peer-reviewed and community owned interoperable data
- Text and data integrated publishing!

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position.

Biodiversity Data Journal 1: e1013 (28 Oct 2013) doi: 10.3897/BDJ.1.e1013











Article info





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Tables and Figures, if present, can be downloaded from the article.

Download all occurrences as Darwin Core Archive

Taxon treatments

Eupolybothrus cavernicolus Komerički & Stoev, 2013, sp. n.

- ZooBank urn:lsid:zoobank.org:act:6F9A6F3C-687A-436A-9497-70596584678C
- SRA project accession http://www.ebi.ac.uk/ena/data/view/ERP00
- ArrayExpress accession attp://www.ebi.ac.uk/arrayexpress/experiments/E-MTAB-1859

prefemur, F - femur, T - tibia, a, m, p stand for spines in respectively, anterior, medial and posterior

- GigaDB http://dx.doi.org/10.5524/100063
- GenBank KF715043
- GenBank KF715049
- GenBank KF715050
- GenBank KF715059
- BOLD dataset http://dx.doi.org/10.5883/DS-EUPCAV
- MorphBank 999021821&tsn=true
- MorphoSource http://morphosource.org/index.php/Detail/SpecimenDetail /Show/specimen id/514

Materials

Download as CSV 🐯

Holotype:

 a. country: Croatia; stateProvince: Knin; locality: NP Krka, village Kistanje, Hydroelectric power plant Miljacka, cave Miljacka II; verbatimElevation: 115 m; verbatimLatitude: 44°00'01.1"N; verbatimLongitude: 16°00'58.5"E; samplingProtocol: hand collected under clay sediment; eventDate: 9 February 2013; individualCount: 1; sex: male; lifeStage: adult; recordedBy: M. Lukić; institutionCode: CBSS; collectionCode: CHP536

Paratypes:

- a. country: Croatia; stateProvince: Knin; locality: NP Krka, village Kistanje, Hydroelectric power plant Miljacka, cave Miljacka II; verbatimElevation: 115 m; verbatimLatitude: 44°00'01.1"N; verbatimLongitude: 16°00'58.5"E; samplingProtocol: hand collected under lump of clay; eventDate: 18 October 2012; individualCount: 1; sex: female; lifeStage: adult; recordedBy: A. Komerički; institutionCode: CBSS; collectionCode: CHP517
- b. country: Croatia; stateProvince: Knin; verbatimLocality: NP Krka, village Kistanje, Hydroelectric

Supplementary material 1

Raw data used for COI delineation of the Eupolybothrus species

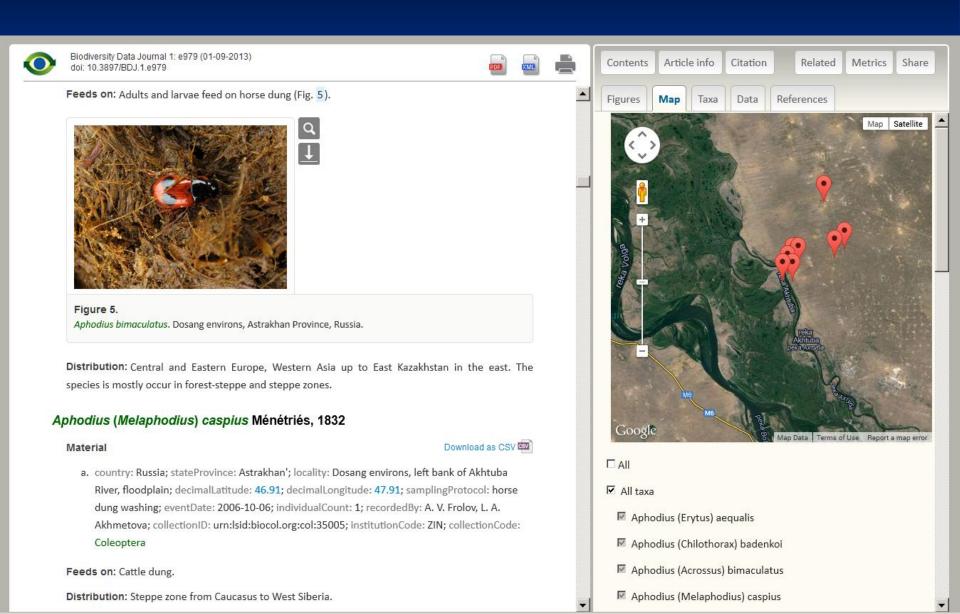
Authors: Stoev et al. 2013

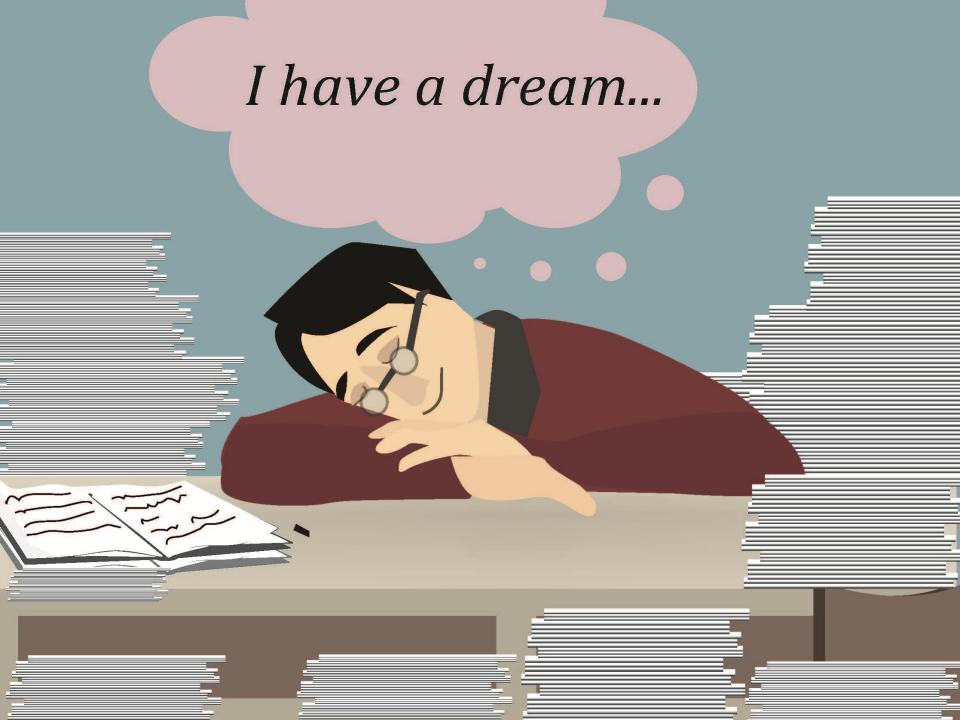
Data type: genomic

Brief description: The archive contains the following data: 1) fasta-Alignment as the basis for all analyses (.FASTA), 2) mega-file for the calculation of the genetic distances and the NJ tree (.MDSX), 3) NJ-tree in Newick format (.NWK), 4) graph of the TCS Software for the Statistical Parsimony method (.GRAPH)

Filename: E_cavernicolus.rar Download file (10.35 kb)

Interactive mapping of localities





Imagine how taxonomic data (if published this way) can be re-used...

... across different domains....



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