

BIG4 field workshop

June 5-11 2016, Havraníky, Czech Republic







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Modern Methods of Systematic Research

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Editorial

Challenges with using names to link digital biodiversity information

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Issues with names as identifiers

- 1. Many names for one taxon (synomyns and other cases)
- 2. Misspelled names
- 3. Same name for multiple taxons (homonyms)
- 4. Chresonyms: scientific name as used by others
- 5. Taxon concept

But

With reclassification, GenBank contained unique canonical name-strings of almost 400,000 (398,740) species and infraspecies of which about 82% could be matched to name-strings in Catalogue of Life. After elimination of known synonyms, 257,702 species name-strings and 20,566 infraspecies matched entries in Catalogue of Life. These represented 13.5% and 1.1%, respectively, of the original name-strings in GenBank and 52.4% and 4.2% of all of the unique canonicalized name-strings.

Definition (*Taxon*)

A set of organisms in nature that form a natural group according to a species concept.

What can a *taxon circumscription* include?

- Textual descriptions of traits that the organisms in the taxon exhibit
- Citations of voucher specimens (types)
- Technical drawings
- Photographs
- High-resolution 3D imaging
- Genetic information
- Comparison to closely related taxa (diff)

See http://zookeys.pensoft.net/about#TaxonomicTreatments

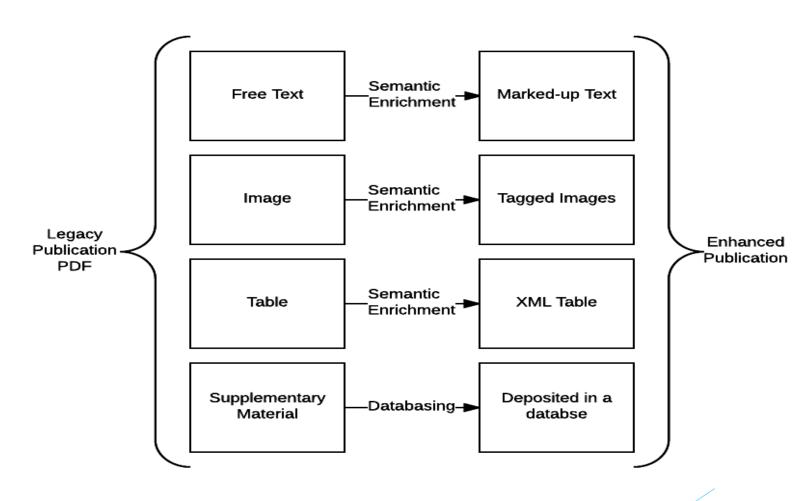
Need for machine-readable data

Since Karl Popper science is moving in the direction of

providing falsifiable hypotheses, i.e.

there must a formally defined algorithm operating on experimental data that could potentially (given the data) disprove the hypothesis. In order to accomplish this, taxonomic data must be made available in machine-readable format, and machine computable format.

Information extraction from a biodiversity publication



Taxonomic impediment

- The Taxonomic Impediment: "worldwide shortage of taxonomists"
- Majority of species undescribed
- ► A lot of "dark taxa": unnamed OTU's

Rod Page, iPhylo blogspot, 12 April 2011

Turbo Taxonomy

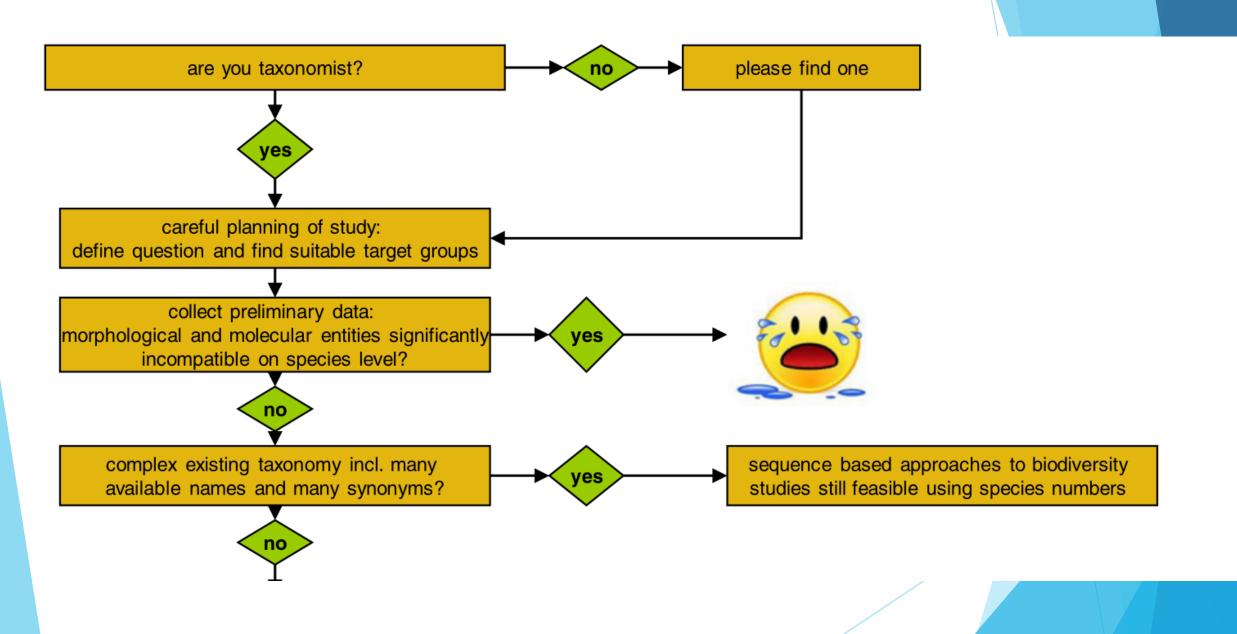
Riedel et al. Frontiers in Zoology 2013, **10**:15 http://www.frontiersinzoology.com/content/10/1/15

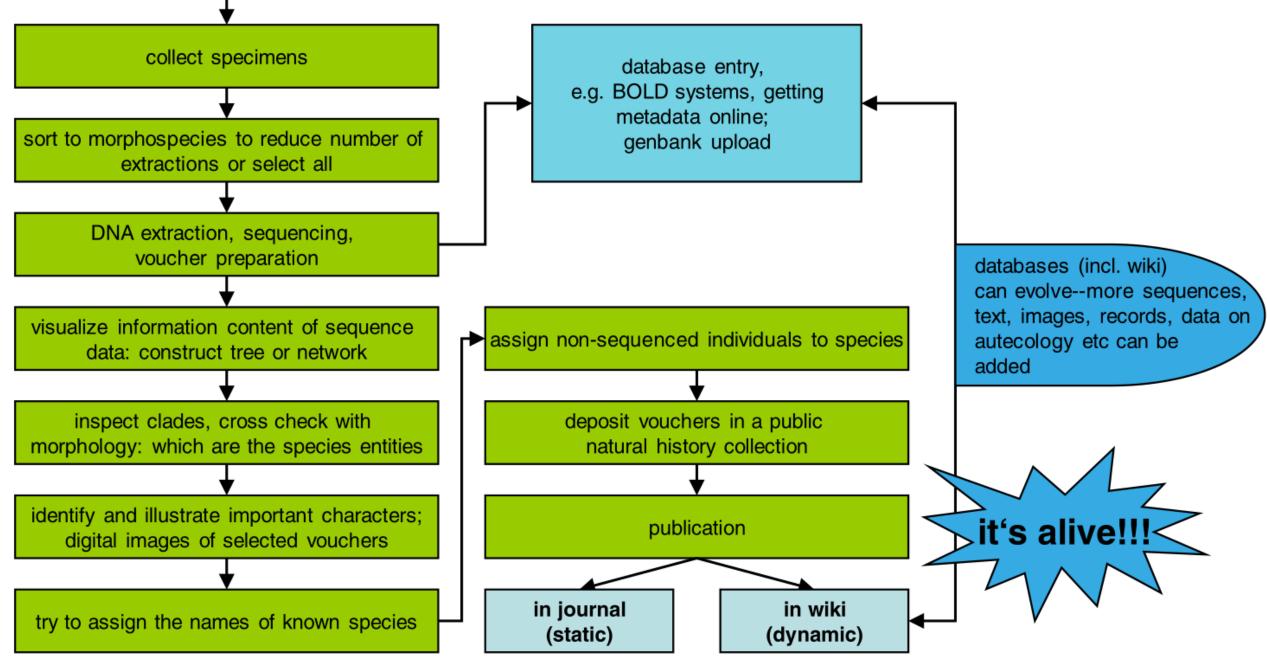


DEBATE Open Access

Integrative taxonomy on the fast track - towards more sustainability in biodiversity research

Alexander Riedel^{1*}, Katayo Sagata², Yayuk R Suhardjono³, Rene Tänzler⁴ and Michael Balke^{4,5}





gure 1 Flow chart of the turbo-taxonomy approach, from project design to publication.

Holistic Taxonomy

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

Komericki & Stoev (2013):

doi: 10.3897/BDJ.1.e1013

Methods of holistic taxonomy

- 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

Slide borrowed from Stoev.



Cybertypes

"Micro-computed tomography: Introducing new dimensions to taxonomy"

Faulwetter et al. (2013)

doi: 10.3897/zookeys.263.4261

"Micro-CTvlab: A web based virtual gallery of biological specimens using X-ray microtomography (micro-CT)."

Keklikoglou et al. (in press). Biodiversity Data Journal.