echinopscis (an extensible notebook for open science on specimens): overview & design principles

TaxonWorks Together (TWT) - 26th October 2023

Nicky Nicolson¹, Eve Lucas²

(1) Digital revolution, RBG Kew, (2) Accelerated taxonomy, RBG Kew



TaxonWorks Together 2023



- Transitioned from software development into research
- Interested in how we can use software development practices in research:
 - Automation
 - Version control
 - Dependency management
 - Continuous integration
- Institutional commitment to use digital technologies to accelerate the process of taxonomy

Community aim: digital extended specimen

- Integrate specimens and associated data across multiple research infrastructures
 - allowing the investigation of wider scale research questions
- How to get there? Activities at a range of scales:
 - Large scale: computational
 - Distributed: lightweight tools, link construction in context by researchers

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- A personal knowledge manager: for creating & linking research notes
- Emphasises linking
- Data stored locally, using open formats
 - Markdown and optional structured data frontmatter
- Works offline
- Extensible architecture plugins for data access and citation processing
- Active user and developer community



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Extend Obsidian for specimen research

- Access of relevant data
 - Specimens (GBIF)
 - Names (International Plant Names Index)
 - Collections (Global Registry of Scientific Collections)
 - People (Bionomia)
 - Literature (crossref)
- Creation of links, spatial and network exploration
- Citation in new work
- Open science working practices

echinopscis.github.io

🥽 echinopscis

Home Team Blog Project -

echinopscis An extensible notebook for open science

"echinopscis" is an experiment in creating an "extensible notebook for open science" - a working environment that allows researchers to write, access data and create links between literature, specimens, names, institutions, people, traits etc.

Key principles:

- Control of your data: as a researcher, you remain in control of your data. The data is stored in text format, on your local machine. Text files are an open format, they will always be accessible without any need for specialised software.
- Open to choose your working practices: we've provided small pieces of functionality that can be combined in many different ways, enabling researchers to be "open to choose" how to organise their work.
- Re-usable skills: any skills necessary to work with this toolkit should be transferable to other open science tools and practices. If you invest in time exploring this prototype software, the things you learn (markdown formatting, bibliography / citation management, document production etc) could also be applied elsewhere in your work, or in other working environments.
- Open science: All code and documentation (and this project site) are managed on github contributions are welcome.

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Welcome

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This is the sample *vault* for "echinopscis" - an experiment in creating an extensible notebook for open science.

Start here

These instructions are a *page* in Obsidian, formatted in Markdown and located on your local computer. You can edit this page - toggle between view and edit mode using Ctrl + E.

By default these instructions are "pinned" so that they will always be visible as we create new pages. Any new pages that you create or open by navigating a link will open in a separate *pane* to the right.

About this demonstration

This vault has been configured with some tools which enable easy access, linking and visualisation of biodiversity informatics data.

The rest of this page gives a worked example of how to use these. Occasionally, some extra information about Obsidian and pointers to the Obsidian documentation has been included in this page in "tip" sections. These look like this - if you click on the coloured bar the contents will unroll.

Tips and extra information about Obsidian >

Functions are available via the command palette in Obsidian - a searchable list of the function 1 backlink D 1944 words 11625 characters

Kev

Arranging specimens: physical working space

- Our physical herbarium includes spaces for layout & comparison of large sets of specimens
- Working digitally means we can access specimen images from anywhere
- But how to effect this kind of interaction in digital environment?



Arranging specimens: virtual prototype





Roadmap

- Personal research environment based on Markdown authoring and linking (now working on "seeding" – pre-populating data)
- 2. Web publication using static site generators (conceptual similarities with the <u>GBIF hosted portal</u> and TaxonPages work)
- **3. Document production**: with structured bibliographic/specimen references
- **4. Dataset production**: mobilisation of content and links into DarwinCore archives for aggregator harvesting



MSc project assessing different options for digital curation, at the Royal Kew Botanic Garden, Edinburgh





Curation of Camellia

Exploring and testing methods of taxonomic curation and identification

of herbarium specimens

Celia Aceae 2023



Thesis submitted in partial fulfilment for the MSc in the Biodiversity and Taxonomy of Plants

Taxonomic Literature (TL-2) – multi-volume work presented in Obsidian KeW

...



Anton Savchenko

@notgaudi

Release alert 👏

I present a digital version of Taxonomic Literature ed. 2, commonly known as TL-2. It can be viewed online or downloaded as an Obsidian database and used offline on you PC. See it here: tl2.io

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Integration: widescale adoption of Reconciliation API

- "strings to things" the echinopscis demo shows multiple workflows where a user translates a piece of text (like a specimen reference) into an entity managed by an external authority (like a specimen record)
- The Open Refine Reconciliation API is now documented as a standard and can be implemented:
 - by the data provider, those that maintain an authoritative dataset
 - Bionomia: <u>https://bionomia.net/reconcile</u>
 - Catalogue of Life: <u>https://github.com/CatalogueOfLife/backend/issues/1265</u>
 - Datasette (generic): <u>https://github.com/drkane/datasette-reconcile</u>
 - by data consumers: developers of taxonomic / data manipulation tools:
 - Open Refine: <u>https://openrefine.org/docs/manual/reconciling</u>
 - Python: <u>https://github.com/jvfe/reconciler</u>
 - Could we build an Obsidian plugin?

Integration: navigation of linked data

- "entity explosion" is a browser plugin that shows links from Wikidata based on the currently loaded page
- Should be possible to develop an Obsidian equivalent, that uses an ID stored in page frontmatter to run a Wikidata query and show useful links



Explore

Background, demo videos, installation information, roadmap and ideas for contributions:

echinopscis.github.io

Source code and documentation:

github.com/echinopscis





nickynicolson



nickynicolson