Nomenclature and TaxonWorks

group notes 2023, 1st Wed of the month

TaxonWorks Docs

Former Digitization Google Doc

<u>Digitization2022 Matrices2022 Nomenclature2022 Open2022 Technical2022 Digitization2023 Matrices2023 Nomenclature2023 Open2023 Technical2023</u>

SUBMIT a GitHub TICKET

6 Dec 2023

SUMBIT a GitHub TICKET

Present: Jim Woolley, Colin Favret, HH,

Updates:

- The TW YT channel now has all our TWT2023 content up, please sharea and subscribe! See https://www.youtube.com/playlist?list=PLPSu6w4rwUI5tRgxboj6V7yxc-GzdOtdw
- Also, the online program https://together.taxonworks.org/2023/ has links to both the videos and presentation PDFs.
- 0.36.0 New reporting and batch functionality

Topics:

Paul Brock - Phasmida Species File (issue No's relate to PSF Issues):

#21 Data issue [part]

Rather confused by the correct process to change data from, say, a syntype to lectotype to show the sequence. Can we tackle *Obrimus bufo* where the LT is now formerly the ST (Oxford) shown here:

https://sfg.taxonworks.org/tasks/accessions/comprehensive?collection_object_id=1560405 Source is Hennemann, 2023: 73. This will then enable me to make other similar changes to the database.

#22 Data issue

A few technical transition issues to solve, but these may take time! One is a typical issue following transition of data, possibly c.300 of these to resolve: *Sipyloidea caeca* translates in TP as:

Descendants and synonyms



- Sipyloidea caeca (Sjöstedt, 1918)

- = Sipyloidea caeca
- = Sipyloidea carterus

'Synonyms' include a 'species' [protonym] with exactly the same name [but no author], which reads 'Sipyloidea caeca (unavailable or invalid, linked to Aruanoidea caeca Sjöstedt, 1918)' in the citations, as well as 'synonym of Sipyloidea caeca'. Not sure if this has to

wait for a merge option, but can anything be done with this to lose the 'unavailable or invalid, linked..' statement? The second 'synonym' [protonym] is an erroneous synonym, which translate in TW as 'Sipyloidea carterus (unavailable or invalid, linked to Aruanoidea caeca Sjöstedt, 1918)' [is this correct in SF for misidentified material or brief synonyms?].

• Other examples include the following – which can include simple change of combination, often for a synonym, sometimes a change in ending of species name:

Descendants and synonyms

=

Prisomera spinicollis Gray, 1835

- = Greenia ornata Brunner von Wattenwyl, 1907
- = Prisomera perlobatum Brunner von Wattenwyl, 1907
- = Lonchodes perlobatus

Descendants and synonyms

=

Paraphasma laterale (Fabricius, 1775)

- = Paraphasma dentatum
- = Stratocles dentatus Piza, 1937
- = Phasma hopii Gray, 1835
- = Olcyphides hopii
- = Phasma lineolatum Audinet-Serville, 1838

MMC

- 1) Any news regarding: the possibility to save monthly permanent archival copies (CoLDP) of the list readily available for the community and access them through a link in TaxonPAges? We discussed the possibility a. to do this with ChecklistBank on monthly releases
 - No work has been done to regularize releases
- b. Zendo publishing approach
 - Nothing done here, need perhaps Debbie to looking into documenting a process.
- 2) TaxonPages: Stats panel:we need to display a new column considering Valid Extant Taxa/ and Valid Fossil Taxa.
 - Issue was added https://github.com/SpeciesFileGroup/taxonpages/issues/197

Colin

How to associate two names for the same person

1 Nov 2023

SUMBIT a GitHub TICKET

Present: HH, Belén, Davide Dal Pos, Holger, Matt, Hernan, Paul Brock, Lesley, José, Anja Erasmus (parasitic arthropods -- microso, SEQ, Delta, museum numbers for specimens, etc. has Specify, and has historical snail data working on georeferencing > has project with SANBI > the "refresh project" snails and freshwater parasites: Anja on marine parasites), Rich, Debbie, Dmitry Dmitriev

Updates:

Topics:

Paul Brock – Phasmida Species File (issue No's relate to PSF Issues):

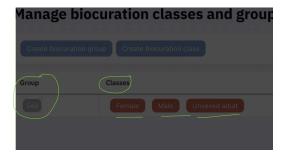
#15 Data issue

Returned to holotype data I keyed on a few weeks ago for *Xerosoma nannospinus*https://sfg taxonpages.github.io/phasmida/#/otus/947347/overview I went to Collection Object and the edit screen but

- cannot see where to enter the sex a female
- I expected the type data to show on the map, but it does not. Not sure what I am missing from the basics here (after perusing the manual), but
- have missing paratype data to add as well



Create biocuration group. Put class members into group as fits.



Then went to Collection Object record to add the sex (female in this case).

And "Brazil" issue posit b/c each of those "Brazil" come from a different place in the database (one comes from verbatim string, the other from the geographic area.



strangeness: see the following:



updated Area and updated cache



Easiest way of doing "paratype"? Do we have "clone" for specimens? Yes, use the "lock" for each value that will be the same, click "Save and New" > add paratype information.

We could think about "lock all"-- discussion seems to suggest folks find it would be useful. Make a feature request y'all if you like this idea.

#11 Combination gueries

- · 2) Phraortes chinensis https://sfg-taxonpages.github.io/phasmida/#/otus/855078/overview the reference to add in Cai p. 27 is Staelonchodes chinensis—this combination is not in TW.
- 3) Ditto Cai p. 28 Ramulus mikado https://sfg_taxonpages.github.io/phasmida/#/otus/933360/overview— I want to add the citation to the synonym listed in Cai, but the reference is to Baculum irregulariterdentata (only Baculum irregulariterdentatum shown as a combination in TW).
 - There are 3 protonyms that we need to collapse down to one.
 - Requires 'merge' fn()

https://sfg.taxonworks.org/tasks/taxon names/filter?per=100¬ specified=true&extend%5B%5D=parent&page=1

- Dryococclus australis https://sfg-taxonpages.github.io/phasmida/#/otus/853177/overview the map for Australia (related to type data of a synonym, probably just a general site) ideally needs removing, as it should just show Lord Howe Island, perhaps Balls Pyramid, if there is a map for that!
- Tropidoderus rhodomus https://sfg-taxonpages.github.io/phasmida/#/otus/853412/overview Philippines needs removing (related to type data of a synonym, thought to be ficticious).

MAPS Issue #15 general discussion

a.

HH:

1. CSF: Anaplecta brachyptera. If I upload images under New Image and select a taxon determination, should I then be able to go to that OTU and see them under the Images tab? Because I can't see them. Not sure what I am doing wrong.

Need GitHub task: organization not showing.

Depictions: "determination" OTUs "ZZZZZ" > New not working

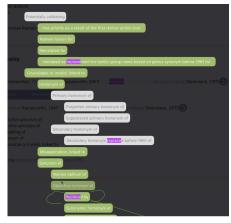
All images directly linked to OTU, should be visible in Browse OTUs

Need to click "apply all" to apply the same metadata

2. DSF: Physogastrinae. Soft validation is incorrect. How to clear?



- b. Heidi saying the "soft validation" assertion that "Objective synonym ..." is incorrect.
- c. DD, should be "subjective synonym" instead (and validation clears)
 - i. (note, when you choose "replaced by" then all statuses nested under that are assumed to be correct).



- 3. Asserted distribution: All the states of India appear to be available except Telangana. Issue?
 - a. https://en.wikipedia.org/wiki/List_of_districts_of_Telangana
 - b. https://github.com/nvkelso/natural-earth-vector/issues/396
 - c. https://github.com/nvkelso/natural-earth-vector/blob/master/CHANGELOG

- 4. Asserted distribution: Is there any way to permanently set our map preference to GBIF, or could it be made the default?
 - a. Incoming issue coming
- 5. PlecSF: Folga luzonica. Some weirdness going on here in the rendering on Edit Taxon Name. Issue?
- 6. CSF: Blattella germanica (and many more). When I search for a valid taxon name, why is there a second copy with no author and a red X? (Chosts? How do I get rid of them?)

Belén - Misspellings and edit taxon name. Recreating the history of a taxon name https://sfg.taxonworks.org/tasks/nomenclature/browse?taxon_name_id=1075173 http://orthoptera.archive.speciesfile.org/Common/basic/Taxa.aspx?TaxonNameID=1120687

4 Oct 2023

SUMBIT a GitHub TICKET

Present: MJY, HH, DDP, Holger, Lily Hart, DD, Jim Wooley, Paul Brock, Holger Braun, MM Cigliano, Belen Cabrera, Michelle Kohler, Carlos Martinez, Tommy McElrath

Updates:

https://together.taxonworks.org/ now with over 120 people registered.

Topics:

HH:.

- 1. Demo from Matt regarding issue with Central African Republic and Plecoptera: Brachypterainae.
 - a. Went through how to debug bad data from bad shapes
- 2. When a synonymized genus is made valid, shouldn't the Manage Synonymy chart come up? It didn't. Is this a bug? (reference: Dermaptera SF: Anisolabella).
 - a. Filed an issue.
- 3. Merge taxa question:
 - a. https://docs.taxonworks.org/guide/Manual/nomenclature.html#how-do-delete-a-duplicate-taxo n-name
- 4. I entered new images in Coleo SF using New Image Task in an effort to generate depiction-ids for the home.md but Depictions button didn't work. Is this a bug? I know this function is working in Plecoptera SF so I didn't know if it was possible for it to not work in another SF. Maybe there is something wrong with my image upload?
 - a. Depiction has to be added to new images using icon at the bottom of image before depiction button will work.

Holger: still unclear: how to fix original combination and rank at nominotypical subspecies: https://github.com/SpeciesFileGroup/taxonworks/issues/3580

- Nominoptypical names must have same spelling to be recognizable in "Fix"
 - Should have same name under parent (at any rank)

DDP: Some info on TaxonPages

MMC: on TaxonPages: regarding the maps problem: we find that only France (French Guiana) and Alaska, Hawaii (USA) are showing problems.

- Discussed 3 aspects
 - o Bad data
 - Bad shapes (our side)
 - o h3 graphs

Paul Brock (Phasmida Species File): Bostranova species - new combination issue. Matt, many thanks for your time in sorting this out during the zoom meeting on 27th. I thought adding the combinations to other Bostranova species would be simple, but I did not make a note of exactly what the procedure was yesterday, as I thought I would remember it i.e. I need to add citations to 25 species, including bifida - this appears as Bostra bifida, rather than Bostranova bifida. You made the change for arcuata and looking at that on-line, it seems I need a combination to appear on the system so the link is correct?

- 1. Can you run through what I need to do again please, so citations show up as Bostranova for the various species, including bifida?
 - a. Pin your source

b.

- 2. The 27th name that did not autotransfer is Bostranova reductedentata, but that is now correct as Phanocloidea (another 2023 reference), but I still need to add the citation, so it appears as Bostranova reductedentata Villet, 2023, within the valid name Phanocloidea reductedentata
 - a. Find the protonym and

Issue with synonymy misinterpreted by software:

https://sfg.taxonworks.org/tasks/nomenclature/browse?taxon_name_id=960337 for Pulchriphyllium bioculatum shows Phyllium magdelainei as a synonym. In old PSF, the valid species for magdelainei is correctly recorded as Pulchriphyllium pulchrifolium. Possibly because of subspecies and changes, the software picked this up as a synonym of bioculatum, in error, although it used to be years ago!

- 1. There seems to be a link missing somewhere, but I need a citation by Seow-Choen, 2018: 675 to read that Phyllium magdelainei is a new synonym of Phyllium (Pulchriphyllium) pulchrifolium (which was in old PSF software). Later, the subgenus was made valid in 2021 by another author, so the correct combination is now, as recorded, Pulchriphyllium pulchrifolium.
- I cannot see this citation at all (under magdelainei in old PSF): Delfosse, Cliquennois, Depraetere & Robillard. 2019. Zoosystema 41(11):211 sciencepress.mnhn.fr >> Note: Type material in MNHN, Paris >> Phyllium (Pulchriphyllium) pulchrifolium
- possibly because Phyllium (Pulchriphyllium) pulchrifolium is not recognised now, although was on old PSF at that time. Presumably I cannot add this back in until issue 1) is resolved.
- 3. Possibly the above is linked to 'unavailable or invalid' citations, linked to Pulchriphyllium pulchrifolium https://sfg.taxonworks.org/tasks/nomenclature/browse?taxon_name_id=960342. Is this likely to be the case here and where I have seen 'unavailable or invalid' citations elsewhere?

I am making progress with keying papers on TW, including new species, new synonym, citations, so getting used to the basics, but this is beyond me - yet!

Tommy McElrath:

How do I update my TaxonPage now? Confused about Setup vs. Main Code.

- Everything through the GH window
- In breakout room

6 Sept 2023

SUMBIT a GitHub TICKET

Present: Debbie, Davide, Jim Woolley, Belén, Holger, Heidi, Lesley, Matt, Dmitry, John Heraty, Rich Flood,

Updates

- Species Files in TaxonWorks
- Potential re"branding" of Wednesday meetings
- Known issues
 - API autocomplete (TaxonPages)
- New report off taxon name filter
- TaxonWorks Together https://together.taxonworks.org

Your topics

Holger and Jim W:

1) Best way to add correct spelling of genus name and make current invalid:

OSF user: "... Agnotecous was originally published with two spellings (Agnothecous on p. 396 and Agnotecous on p. 406). This mixed usage remains in the Mélanges Orthoptérologiques reprint without any errata. The index to the latter work itself uses Agnothecous. Saussure later (in Biologia Centrali-Americana 1:256) used Agnothecous, which is treated as a first reviser action under ICZN art. 24.2.4. Additionally, Neave explicitly selected Agnothecous in 1939 (in Nomenclator Zoologicus 1:91). I believe either of these alone would make Agnothecous the correct spelling for this genus."

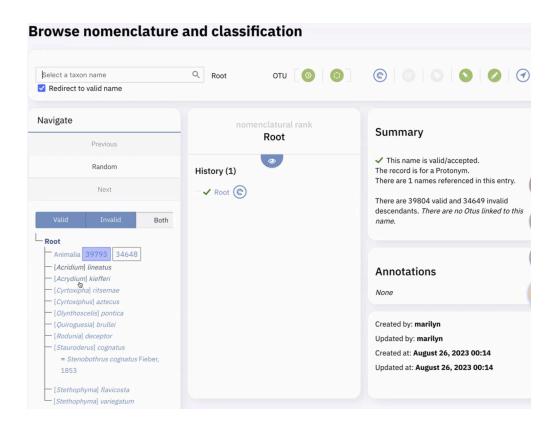
simply add Agnothecous (missing in OSF) – make Agnotecous a synonym of it – verify citations at genus and species?

- Create the correct spelling
- Synonymize the misspelling to the correct spelling
- At this point you can move the synonyms as needed
- This will preserve citations as need

JW has family-group names issue (parallel), 2 of them.

- will follow steps above, citing person and publication who noted / asserted correct spelling
- is there a choice? yes, 'incorrrect original spelling'
- For family group names "autocorrect"
- 2) Names under ROOT: cannot find old GitHub issue on how to fix this (10 unsuccessfully migrated combinations)

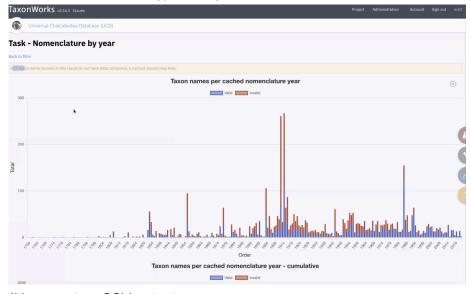
Turns out: these seem to be protonyms not combinations, they will need checking.



- set the parent or fill out data
- 3) Taxon numbers by year? (nomenclature stats)

Matt showing Unified Filter (and new function added: e. g.

- Filter nomenclature,
- send result to chronology stats, gets



- JH requests a CSV output
- MJY could also output a SVG version of the graphic too, if desired.
- To JH, et al see samples of what's possible https://www.chartjs.org/

4) Heidi: (TaxonPages)

- 1. Config: Punctuation in config files?
- 2. Project: What to put under citation?
- 3. Metadata: Are initial caps critical under search terms?
- 4. On Home page: how to configure link text under search category?
- 5. Repository for Isoptera Catalog?
- 5) Belén: (Digitization) Just if we have time!!!
 - 1) The order of the geographic areas from coordinates is not hierarchical. E.g. Congo
- 2) We would like to have the shape corresponding to georeferenced points at the level of species. So if we have a species in taxon pages with only coordinates the map can also have a painted area for the general distribution.
- 3) Remainder: Hard validation on assigning type status in comprehensive task. E.g. Assigning paratype status twice for the same collection object.

JH: question, download PNG

DD: taxonpages

- etymology panel
- part of speech panel
- tiff file issue

2 Aug 2023

SUMBIT a GitHub TICKET

Present: Debbie, Dmitry, María Marta, Belén, Holger, Daniele, Roger, Heidi, Oliver, Dash [You], ...

Updates:

- TaxonWorks Together 24 26 2023, please submit ideas now!
- Companion software **TaxonPages** live later in August. Public website software pulling data from your TaxonWorks Project databases.

Topics:

MMC: On adding images on an existing species that is currently used in a different combination from the genus in which it was originally described: In the TW doc

(https://docs.taxonworks.org/guide/Manual/Sources/#add-more-species-information) it is recommended to use the OTU attached to the combination. It is mentioned that "this is an area of the database that is being worked on, but this should facilitate downstream display of the new information'. Would this still be the best procedure?

DP: MMC, that section of the manual comes from UCD folks, so indeed, needs revisiting. Dmitry (DD) will be joining us and can hopefully answer this question.

DD: From 3i point-of-view, different from Matt and UCD procedure is to attach images to protonyms, not ... combinations

- prefers to keep images, host info all attached to single OTU
- whereas UCD wants to keep ^^^ linked to exact name which it is listed in the paper.

DDP: How to handle citation of subspecies when their status changes.

species > because now a subspecies but had entered all of it as species, with some info then added at subspecies level when he realized it was a subspecies.

DD "cheats" -- changes the species entry to subspecies > adds species again > updates parent of the subspecies (OTU). Minimal changes to the database.

OK: species described in genus A, at some point (50 years later), catalog now shows species in 2 genera B and C.

Daniele, Hoverflies, NW of Italy region in TW. Increase distribution information for North of Italy. An interactive key for CitSci for hoverflies. TW a solution for these purposes.

DPeters: Are OTUs the right way to represent unpublished names? How to store this as more than just a text field?

DD: yes, attaches OTU to higher rank. (TaxonID and rank). No citation, b/c it's unpublished. (DD puts citation on TaxonName).

DD to DPeters: Create OTU w/o a taxon name.

Create OTU, put name on it, for specimen data upload. OTU doesn't have to be attached to TaxonName.

Roger (RAB): I plan to be on late-ish, maybe trying again in open forum: want to talk about an FDA multi-entry key project.

OK: Digitization, using "Save and New" took 30 to 60 seconds instead of being very fast. to OK: please ping us on Gitter when this happens, so we can help in real-time to determine if it's on our end or yours.

DPeters: when does sandwich get updated (to try out new code, please)? to Dash: please ping Hernan on gitter.

RB: "remote entry key" for FDA.

- no google docs on FDA computer.
- have to do DEV at home, using pull requests.

5 July 2023

SUMBIT a GitHub TICKET

Present: [Your Name Here], Debbie, Matt, Jiri Hulcr, Miranda Barnes, Lesley, Holger, ...

Updates:

TaxonWorks Together 2023

24 - 26 October

Release soon (Friday?) with cached map software. IF you want to help, please test on sandboxes

- create georefs or asserted distributions
- will labels / maps be in English? (Jose found another map tile set, from GBIF, but it's slow)
 - still an open issue.

Your Topics:

Jiri Hulcr - Doing a co-revision (beetles and fungus), connected ecologically

- we handle multiple codes of nomenclature / including fungi? yes
- otu otu, otu specimen, specimen-specimen
- filter nomenclature, filter spatially, filter by "in project"
- some parts of revision pieces can be built in TW (like "Paper Catalog")
 - another group is building a LaTeX version
- observation matrices ... descriptions can be written in TW
- "specimens examined" handling coming to TW
- lists of biological associations
- GloBI integration, Jorrit Poelen is pulling associations data now from TaxonWorks
- Using TDWG media standard Audiovisual Core Multimedia Resources Metadata Schema

-

JH: Graph database question? Where do folks begin? to learn how to manage graph data? MJY: TW is in PostgreSQL, a relational database format.

what are you asserting versus inferring

JBW: I am having trouble cutting and pasting from a pdf. Maybe someone can help?

7June 2023

SUMBIT a GitHub TICKET

Present: [Your Name Here] Matt Yoder, Debbie Paul, Holger, Lesley Deem, Valeria Trivellone, Chris Dietrich, Jose, Rich, Belen, Geoff, Heidi, Guy, Dmitry, Valerie T,

Updates:

Your Topics:

JBW: this one is actually nomenclature appropriate:

1- Here we go again. I am trying to set up the following relationships in UCD

Chalcididae Latreille

Phasgonophorinae Steffan (treated as subfamily by Cruaud et al. 2021)
Phasgonophorini Steffan [this is the part I can't seem to do]
Phasgonophora Westwood

DP: mischief managed ... the issue was that there was a child at the rank of tribe, and when changing the parent of that name to tribe, the software was unhappy. Fixing the rank of the child, then it was possible to have the parent at the desired rank.

Issue, lost the original spelling of the group ... needs to attach to the tribe (not sub-family)

MMC: this is on TaxonPages

1- Stats: As a user I would like to see the statistics for each Taxon showing the number of descendants taxa for all categories. For example for Subfamily Melanoplinae, below. Would it be possible to show also the number of Tribes? Besides, it shows results for Families?

Taxa: Species: 1575; Genera: 150; Families: 12 Names: Species: 1906; Genera: 200; Families: 23

Holger (addition to the above): Taxa should probably be number of valid taxa, and Names number of protonyms (e.g. on page of genus: number of valid species, and protonyms of species and genera, no combinations of species names)

Example genus *Diponthus*: Taxa: species: 25; genera: 1 Names: species: 33; genera: 1

There are in fact 21 valid species and altogether 26 specific protonyms.

GENERAL / Other topic questions are also welcome. Please put them there ...

Valeria's and my (Chris) new project focuses on documenting associations between insect herbivores (Hemiptera), plant-pathogenic (and other) bacteria, and vascular plants. Valeria already has a 3i database documenting known tri-tropic associations between these three lineages (kingdoms): http://trivellone.speciesfile.org/index.asp

We would like to have a project in TW that enables us to represent these kinds of biotic associations and incorporates the full taxonomic hierarchy for each OTU/species for all 3 groups (insects, plants, bacteria). What is the best way to do this. Ultimately we also want to be able to visualize/analyze associations at different levels in the taxonomic hierarchy as well as among clades on current phylogenies for each group.

Challenges

- Curating data
- Computing
- Visualizing data
 - Discover

Data

- Phylogeny
 - Requires OTU ids to swap labels
 - 4 Way Phylogenies
 - Cube this

Constraints

We don't want to update nomenclature

Existing

A 3i project with known associations already presented.

Needs

- Let me see at genus level
- Select different hierarchical levels
 - evaluate the pattern at different level

- Hierarchical search
 - o How many genera, etc.
- "Phylogenetic dimension"

DP to Chris and Valeria? Which Plant Nomenclature / Bacterial nomenclature resource would you point to if you had the option?

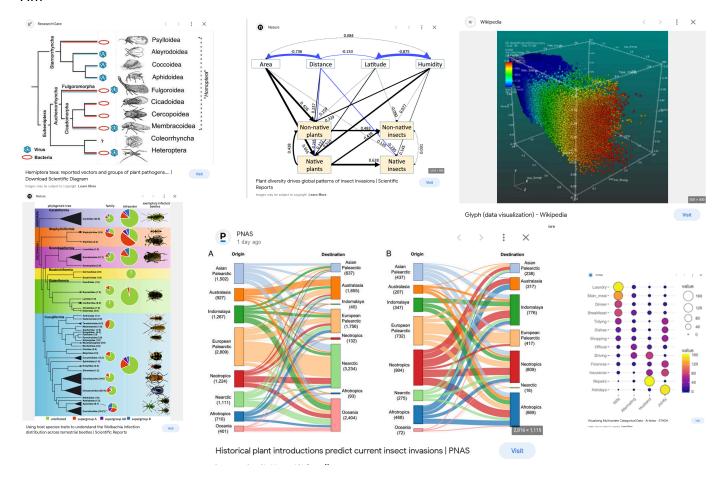
Wants to visualize the relationships between these groups, at various taxonomic rank levels to discover patterns of associations.

3i covers most of the species that vectors of these bacteria that are plant pathogens. Valeria has a pathogen database.

There are databases of plants (CoL, IPNI?, ITIS?, ...)

VT: Biotic Associations as a separate project in TW. Want to uncover / visualize at a diff hierarchical level. See GloBI

Hm



3 May 2023

SUMBIT a GitHub TICKET

Present: Davide, Belén, Holger, Martina, Debbie, ... Your Name Here?, ... Dmitry, Matt, Heidi, Jim, Lesley, Martina, Geoff, ...

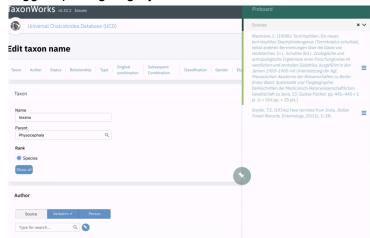
Updates:

- Some key features to remind users about in Nomenclature
 - Yellow facets

Your Topics:

Heidi

- Under the author section of New Taxon Name page could we have a "clone last source" button, or a list
 of the previous 3 or 4 sources. This would be hugely helpful when entering many species from the
 same paper.
 - o easy to forget pinboard is there b/c if nothing is pinned, you don't see it.
 - suggest putting it "grayed out"



- thanks Belén for adding an issue for this!
- Somewhere on the same page, could there be a little checkbox to indicate when a species is the (type
 of the genus) genotype? And could this autopopulate that info on the genus page? This would be super
 wonderful!
 - We illustrated a possibility of creating a new species in the context of a new genus
 - Make a button or checkbox please for this?
 - Consideration:
 - Maybe Taxon Names "Quick Forms?"
- Could we have a button on the Navigate tree (left of screen in nomenclature) to expand-tree so that we can see the entire thing at once? Also would be hugely helpful.
 - o need for building a taxonomic history, where are you in this process, needing to track progress
 - o ability to lock in a view context (lock genus node in place)
 - concept chrono browser

JBW:

1- Here's one from Natalie:

0

Hi all

I am uploading a lot of images from a Mymaridae publication, generally more than one image per OTU: once I have bulk uploaded them to the OTU, and working from the "depictions" screen (as below), I add the label / caption (from the edit button) + the citation (from the radial). I find it very awkward to know which images have already been updated and

which remain to be done - is there a better way of doing this? I have about 100 images to upload and am being driven slowly but surely mad...

- Let's explore:
 - o Image filter -> Per OTU -> right click to work in that context
 - Potential for addition of Images in depictions without label/caption facet
- Let's explore CSV

And also: is there a way of seeing the name of the images directly (i.e. without opening the "inspect" window)? (I have renamed all the images with page number + taxon, e.g. "p.61_figs- 1–3_Acmopolynema" so seeing the image names would make it much easier to check they have been attached to the correct OTU).

Any help appreciated (this can of course wait until the next TW BBQ – I am on leave next week so hope to temporarily regain my sanity \bigcirc).

Have a fantastic weekend everyone!

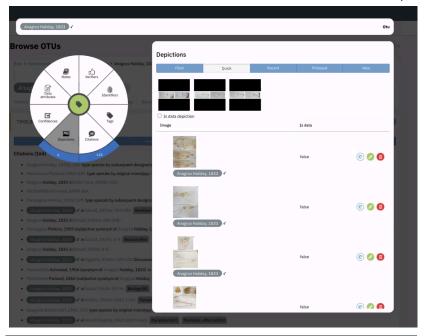
r

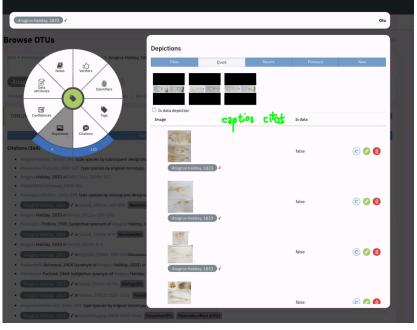
TICKET Made: Filter Images, w/ w/o depictions not working as expected / needed. False positives https://github.com/SpeciesFileGroup/taxonworks/issues/3395

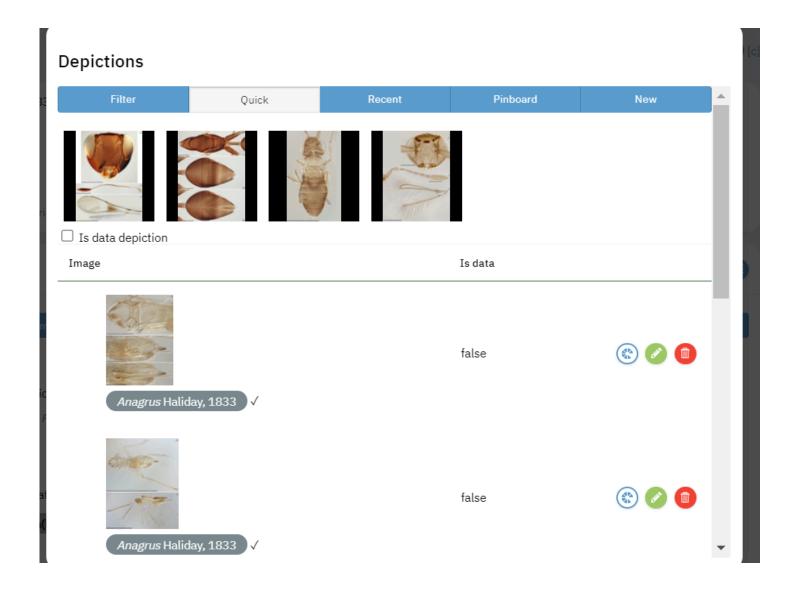
- hard to remember, can't see / find tell
 - which have captions added
 - which have citations added

GitHub Tickie Made: UI Change Requested https://github.com/SpeciesFileGroup/taxonworks/issues/3396

- add columns to this view to show "has citation" or "has caption"







Holger, question for Dmitry:

In a new paper in Zoologica Scripta (Early View) a new cave cricket species is mentioned in the paper as such, but almost all nomenclatural information (link with ZooBank LSID, diagnosis & description, holotype with locality, etymology) is only mentioned in a simple Word document available under Supporting Information. Is this valid? The depository is mentioned in the paper itself.

DD: Use the following checklist in this case

- No, the electronic only publication can be only in PDF format
- The separate file must be treated as its own document, it does not "inherit" metadata from the "parent" publication
- The data of publication must be *inside* the document
- The new document must have all the typical information required (e.g. holotype repository) for a name to be made available

Images of the same thing -> match by MD5

DDP:

I have a question on *incertae sedis* soft validation. (What was the answer here? "not fixable?" in Soft validation?)

 UI improvement coming, and explanation (in this edge case not enough data present for a fix to be automatically applied)

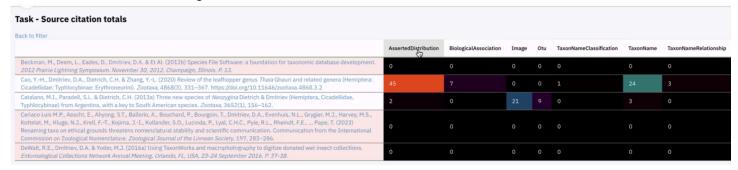
Belén:

In Filter People, Full Name field, I would love to have a gray text to clarify that the only way to put the name so I get results is entering first the last name and then the first name. I can learn this practice, but I learned it by asking... Maybe if there is a clue there the new people can save that time.

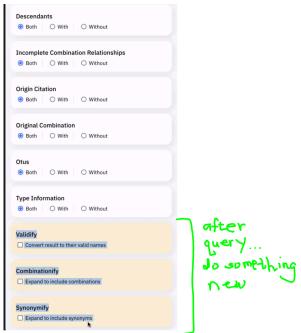
Many of the journals I work with list the names of the authors using the first name first and then the last name. It's not clear for me that we have to use a particular order, unless somehow someone makes it clear to me. https://github.com/SpeciesFileGroup/taxonworks/issues/3394

clue being added in the form of grayed-out text in the search box that helps people know what to enter
in the search in terms of name order.

TOOL / FEATURE Hint: Using "Linker" from Filter Sources to see what's done or not done.



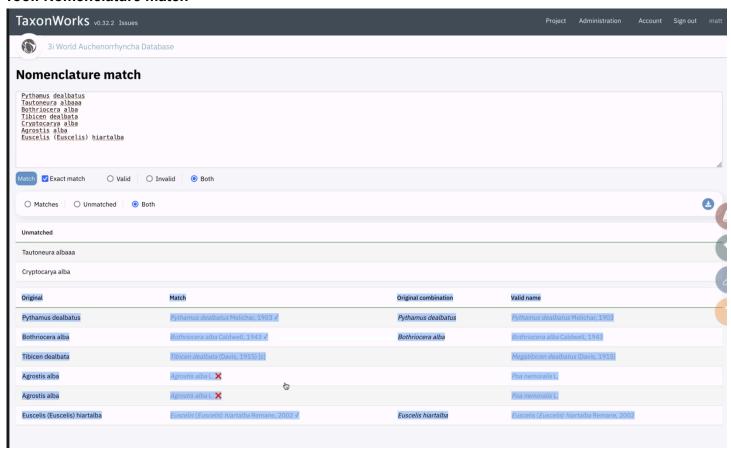
HINT after Filter Sources, new options become available to filter on ... in left column



Combinationify: "Expand my results to any where one of my names have been used in a combination" all names all combinations

Synonymify: expand current results to include all synonyms

Tool: Nomenclature match



5 April 2023

SUMBIT a GitHub TICKET

Present: Matt, Jim, Heidi, José, Hernán Moderator: Jim Woolley (thanks Jim!)

(Regrets: Debbie, Matt, Dmitry as we are at / hosting the Catalogue Of Life Global Team Meeting)

Updates:

Matt demonstrated the New Biological Associations task

Your Topics:

1 March 2023

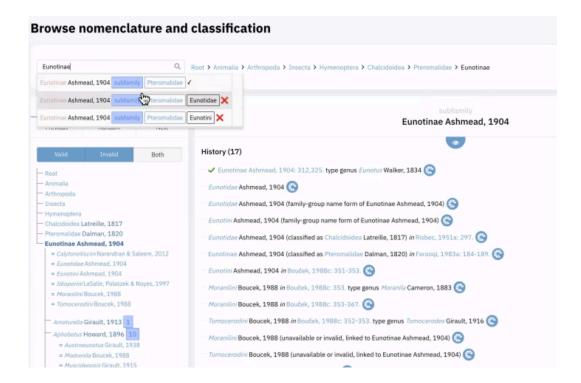
SUMBIT a GitHub TICKET

Present:

Updates:

Your Topics:

I have an issue with interpretation of Eunotidae: John



DD and MJY to write up what to do / how to handle this case ^^^.

Holger and Belén:

☐ How to create a new combination which is a variety (to add or correct citations to old references). It creates a subspecies name instead of the correct one that is a variety. Example case: Stictogryllacris submutica var. neavei

http://orthoptera.speciesfile.org/Common/basic/Taxa.aspx?TaxonNameID=1131625

please look here
 https://github.com/SpeciesFileGroup/taxonworks_doc/blob/archive/manuals/NOMENCLATURE_BASICS.md and note this information will soon be at docs.taxonworks.org

DD: I add those as subspecies, since varieties are outside of the scope of ICZN and varieties should be treated as subspecies. That is why variety could be entered only for original combination, but not for subsequent combination. If needed a comment could be added with the radial annotator.

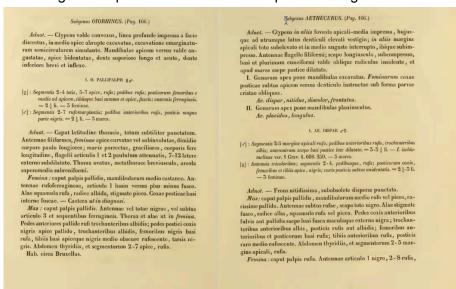
The visible string can be over-ridden by using the verbatim name of a combination.

☐ In our data most of the taxon names that were originally described as a variety are considered invalid. They have (excluded: infrasubspecific) as status. How could we fix this without adding the "Valid" status to force their validity?

DD: delete 'infrasubspecific' status where it is not appropriate, it should only be applied to the names described after 1961. You can use Filter nomenclature to find all taxa which have the status and described before 1961.

Davide:

I have a general question about the description of subgenera that are now genera.



All the sp described here ^^^, now don't have the author above in parentheses.

DD: sp author should be in parentheses, if for example, sub-genera moved to genus. If the genus name portion changed, you need parenthese.

if species moved ... then author not in parentheses.

1 February 2023

SUMBIT a GitHub TICKET

Present: Matt Yoder, Cam Webb, Heidi Hopkins, Jim Wooley, Geoff Ower, Roger Burks, José Lui, Maria Marta, Davide DP, Samson, Lesley, Rich (Dmitry D will be late)

Updates:

• Will need to adjust to a new Gitter / or replacement for Gitter (after Feb 6th)

Your Topics:

JBW (I may be late joining the meeting as it is a BBQ day)

1- I can't seem to update the taxonomic history (subsequent combinations) of *Closterocerus purpureus* (Howard), p. 71 in source 127628

4 January 2023

SUMBIT a GitHub TICKET

Present: Jim Woolley, DD, HH, MY, IG, LD, JH

Updates:

- Verbatim source rendering bug should be fixed today
- Work on unified filters is ongoing and making nice progress
- Year theme "integration", getting taxonworks to talk to other resources.

Your Topics:

JBW

- 1- nomen nudum, source *****, how much information to curate, what do I tell the authors?
 - Holotype deposition

Dmitry

- In Code 5 there is a good chance they will reduce requirements
 - This means that retroactively there may be some names that are valid then that are not now.
 - "Permits" well known (an not so well known) specialists to make the odd mistake in publication without "punishing' them (e.g. JBW's issue this week)

Davide

- Up to 1500 names in ich
- Some new people seeing potential to store images
- Doing some work in the next couple months to recruit others
- !! Need to check back at his email requests re morphological matrices

10°27'N, 84°43'W, 26.ii.2003, J.S. Noyes" (MZLU, NHMUK).

Etymology. From the Latin *pumilus* = dwarfish, little, referring to the small size of this species.

Closterocerus purpureus (Howard) comb. rev.

(Fig. 199)

Necremnus purpureus Howard, 1897:164-165. Holotype female, type locality: West Indies, Grenada, NHMUK, examined.

Closterocerus purpureus (Howard); Waterston (1915:330).

history not in TW

Chrysonotomyia purpurea (Howard): Bouček (1977:7).

Chrysonotomyia (Achrysocharella) purpurea (Howard); De Santis (1983:22). Chrysonotomyia purpureus (Howard); LaSalle & Schauff (1992:9).

This species belongs to *Closterocerus* s.str., but it is difficult to establish its species identity because the head, including the antennae, are missing. Especially the antennae are very important for the identification of *Closterocerus* species, but to a certain extent also the head, e.g. the sculpture on, and size of vertex, are used to separate species. Therefore it cannot be included in the key.

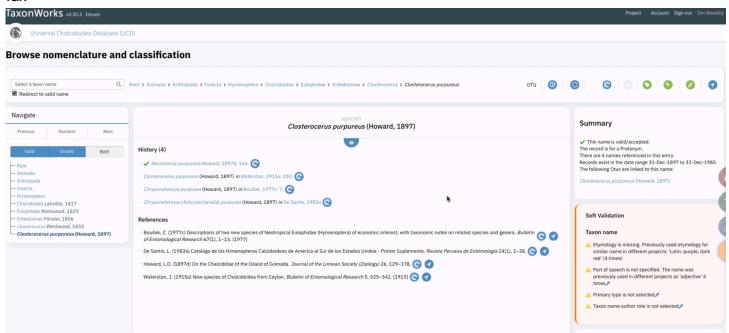
The holotype female (Fig. 199) lacks a carina along anterior margin of pronotum. Mesoscutum and mesoscutellum have quite strong reticulation; mesoscutum is metallic purple and mesoscutellum metallic purple with median part golden-green; midlobe of mesoscutum has short setae of an uncertain number. Legs: coxae difficult to see; fore and mid femora predominantly white, hind femur dark brown with apex white; tibiae white; tarsi white. Fore wing hyaline with parts below stigmal vein and along apical margin infuscate; speculum is closed. This combination of features does not match any of the included species and I therefore keep *C. purpureus* as a valid species of *Closterocerus*.

Hosts. Unknown.

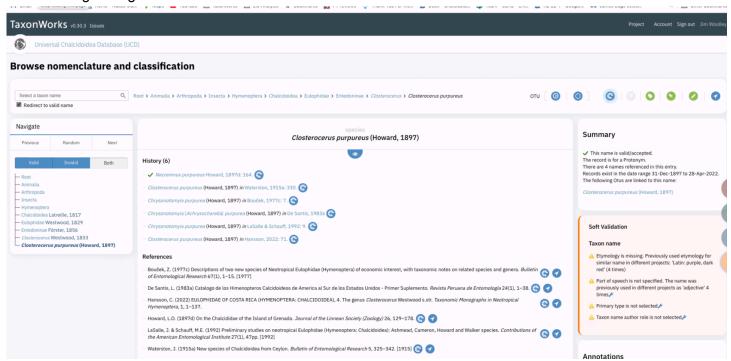
Distribution. Grenada (Howard 1897).

Start:

Tax



After working through it:



Cam Webb, with Stefanie Ickert-Bond, on Ephedra

- 20 years of data
 - Large set of specimen records
 - Large library of micromorphology
 - Field photography
 - Specimens

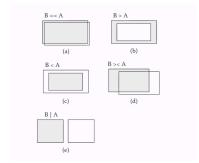
- Sequencing data
- Taxon Concepts efforts
- Goal: to enable the next generation monograph (cf. Zapata and Daly NSF workshop)

Question: are folks using TW to making / building monographs?

- paper catalog
- Dmitry Mozzherin
- About the Dynamic Monograph: see https://ssbbulletin.org/index.php/bssb/issue/view/299
- TW Public Pages coming by end of 2023 >

Cam background

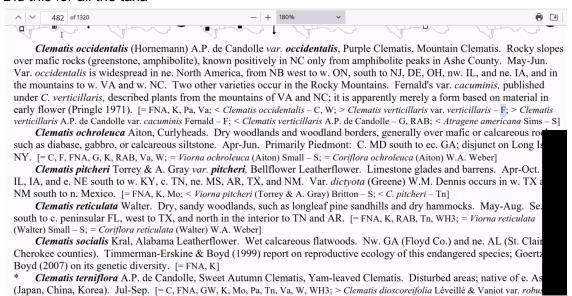
- ABI managing TaxonConcepts grant (https://alaskaflora.org/)
 - Practical way of manage this
 - Encode this
- Background:
 - o A name on its own doesn't specify a specific set of organisms
 - The set of individuals can change a great amount over time.
- TaxonConcept Name + Publication == 'secundum"
 - A "diad of information"
 - o Similar to a circumscription, but the latter has a set of data that is also linked in
 - "The idea the taxonomists had in their head
- Domain
 - o Flora of Alaska
 - 50-60 years of data
 - Names have had very different meanings
 - Goal is to connect the histories of the concepts through time



- 5 ways that hey can relate
- "Superproperty" -> intersect ion of the 4 top properties
 - You can estimate, but can't tease out the precise nature of the concepts
- E.g. Alen Weekly biggest example using these concepts via the Flora of the NorthEast(?) project

https://ncbg.unc.edu/2022/04/26/new-edition-released-flora-of-the-southeastern-u-s/ (perhaps)

Did this for all the taxa

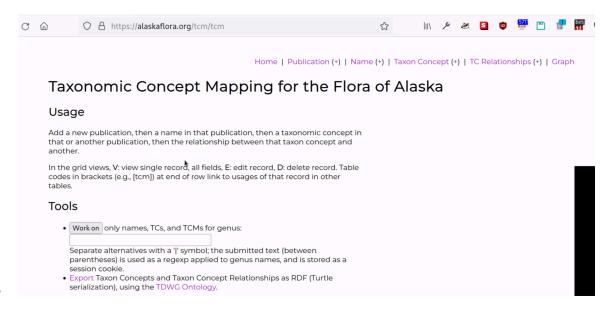


Rendering wise:

early flower (Pringle 1971). [= FNA, K, Pa, Va; < Clematis occidentalis – C, W; > Clematis verticillaris verticillaris A.P. de Candolle var. cacuminis Fernald – F; < Clematis verticillaris A.P. de Candolle – G, R.A.

TCM Script from Cam

- https://github.com/camwebb/tcm
- https://camwebb.info/files/webb2021_tcm.pdf



С

Taxon Concept

Code	Name	Publication	BHL pg1	
R. abortivus sec. Co	Ranunculus abortivus L.	Cody, Yukon Flora		V E D [tcm] [tcm]
R. abortivus sec. FN	Ranunculus abortivus L.	Whittermore FNA Ranunculus		V E D [tcm] [tcm]
R. abortivus sec. Hu	Ranunculus abortivus L.	Hultén, Flora AK Book		V E D [tcm] [tcm]
R. abortivus sec.	Ranunculus abortivus L.	PAF		V E D [tcm] [tcm]
R. abortivus sec. We	Ranunculus abortivus L.	Welsh 1974		V E D [tcm] [tcm]
R. acris sec. FNA	Ranunculus acris L.	Whittermore FNA Ranunculus		V E D [tcm] [tcm]
R. acris sec. Hulten	Ranunculus acris L.	Hultén, Flora AK Book		V E D [tcm] [tcm]
R. acris ssp. acris	Ranunculus acris ssp. acris	PAF		V E D [tcm] [tcm]
R. acris var. acris	Ranunculus acris var. acris	Welsh 1974		V E D [tcm] [tcm]
R. acris var. frigid	Ranunculus acris var. frigidus Regel	Welsh 1974		V E D [tcm] [tcm]
R. aquatilis var. aq	Ranunculus aquatilis var. aquatilis	Whittermore FNA Ranunculus		V E D [tcm] [tcm]

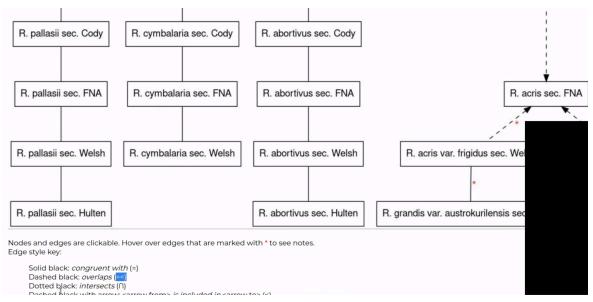
Taxon Relationships

Home | Publication (+) | Name (+) | Taxon Concept (+) | TC Relationships (+) | Graph

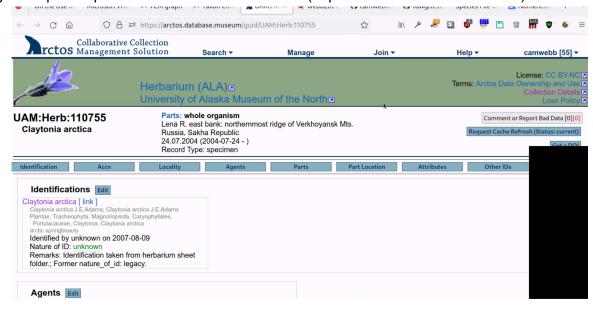
TC Relationships

TC1 newer	Relation	Synonymy	TC2 older	Publication	Graph		
C. pallasii sec. PAF	is congruent with		R. pallasii sec. Cody	inferred by Cook	1	V E	
H. cymbalaria sec. PAF	is congruent with		R. cymbalaria sec. Cody	inferred by Cook	1	V E	
R. abortivus sec. Cody	is congruent with	k	R. abortivus sec. FNA	inferred by Cook	1	V E	
R. abortivus sec. FNA	is congruent with		R. abortivus sec. Welsh	inferred by Cook	1	V E	
R. abortivus sec. PAF	is congruent with		R. abortivus sec. Cody	inferred by Cook	1	V E	[
R. abortivus sec. Welsh	is congruent with		R. abortivus sec. Hulten	inferred by Cook	1	V E	[
R. acris sec. FNA	includes		R. acris var. frigidus sec. Welsh	inferred by Cook	1	V E	
R. acris sec. FNA	includes		R. acris var. acris sec. Welsh	inferred by Cook	1	V E	[
R. acris ssp. acris sec. PAF	is included in		R. acris sec. FNA	inferred by Cook	1	V E	
R. acris var. acris sec. Welsh	is congruent with		R. acris sec. Hulten	inferred by Cook	1	V E	[
D paris var frigidus sas Walsh	is congruent	1	R. grandis var. austrokurilensis	informed by Cook	١,], ₋	

Visualization of the relationships



- visualizing possible gaps
- teaching?
- overlapping with range maps?
- discovering errors in your data or relationships?
- overlapping BHL data
- each concept assoc with specimens, they can be linked out
 - then make inferences if specimens need annotation / revision
 - agencies have different master lists ... they follow one set of names ... when it turns out is much more complex. They do "niche modeling" based on name usage, but they may be lumping totally diff things
- linking concepts to specimen identifications in Arctos (https://arctos.database.museum)



- Citations vs. Housekeeping
 - o for Publication
- Synonym encoded in the same table?
 - o Do you mean in the same form, but in a different place?
- Graph visualization
 - o How is it used?
 - Per citations pivots
- Each concept has specimens
 - Visualizations
 - color indicate overall data presence
 - what has specimens
 - what has original description papers
 - heatmaps of data coverage, concept
 - minimum information
 - !!! Agency master lists have to be mapped !!!
- "Agency to Agencies"
- Agency -

students in SySBio course, using the visualizations, to learn how names work and no one name list works, and names will change.

- MJY bringing up Bertram Ludaescher's 21-relationship (geospatial relationships RCC5, RCC20 or 21), has Cam investigated these?
 - \circ just a bit, but thinks the 5 relationships are good enough. (A == B, A > B, B > A, etc)
 - MJY: talking with Hong Cui, re SKOS concepts (like "same as") turns out not to be helpful in computing.
- TDWG TC(transfer)Schema
 - 5 years of of modernizing TCS ongoing https://github.com/tdwg/tcs2
 - encodes
- reasoning EulerX
 - o infer all other relationships
- Steffi's data in RDF? Yes, from the same tool.

```
t:accordingToString "Cody, M.J., 2000, Flora of the Yukon Territory, NRC Research Press, Ottawa".

1766 a tc:TaxonConcept; rdfs:label "E. acris ssp. politus sec. Hulten"; tc:nameString "Erigeron Acris ssp. politus sec. S. & K."; tc:nameString "Engeron Acris ssp. politus sec. S. & K."; tc:nameString "Engeron Acris ssp. politus sec. S. & K."; tc:nameString "Engeron Acris ssp. politus sec. S. & K."; tc:nameString "Engeron Acris ssp. politus sec. S. & K."; tc:nameString "Hans Schinz and Robert Keller, 1909, Erigeron Linaeus, Flora der Schweiz, Teil: Exkursionsflora, 3, 577, Albert Raustein. Formerly Meyer & Zellers Ver 1765 a tc:TaxonConcept; rdfs:label "E. acris ssp. politus sec. PAF"; tc:nameString "Engeron acris ssp. politus (Fr.) H. Lindb."; tc:accordingToString "Elven, R. and Murray, D.F. and Razzhivin, V.Y. and Yurtsev, B.A., 2020, Annotated Checklist of the Panarctic Flora (PAF) Vascular pl. 17667 a tc:TaxonConcept; rdfs:label "E. acris var. asteroides sec. Porsild"; tc:nameString "Engeron acris ssp. asteroides (Andrz.) DC."; tc:nameString "Engeron acris ssp. asteroides (Andrz.) DC."; tc:nameString "A. E. Porsild & William J. Cody, 1980, Vascular Plants of Continental Northwest Territories, Canada, National Museum of Natural Scien 17671 a tc:TaxonConcept; rdfs:label "E. acris ssp. debilis sec. Cody"; tc:nameString "Engeron acris ssp. Piper 1986"; tc:nameString "E
```

Next.

- talk with Steffi, about short-comings perhaps, for this existing project.
- we have had plant groups investigate using TW, but didn't choose yet
- we did make some changes to our database to accommodate Plant data
- then, what do we do to actually build this out, working with Jose,
- what is your timeline Steffi? (after Feb, start writing a grant to NSF Arts program)
 - does it make monography easier?
 - wants ways as an attractive product at the end?
 - can people explore, to do NM in TW?
- would this be compelling to NSF? to put this work into TW? current work is proof-of-concept
 - helping agencies, helping students ... would be compelling I think (DP)
- moving legacy data into TW is a challenge and takes time
- existence of a real dataset behind-the-scenes is a strong point for MJY
- what next?
 - meet after February, to discuss more about what the Grant you want to write will include, could include with regard to TW and TW Public Pages functions

MJY: sharing about Unified Filters software