

Data Aggregator – Tutorial Session 1 Lesson 1

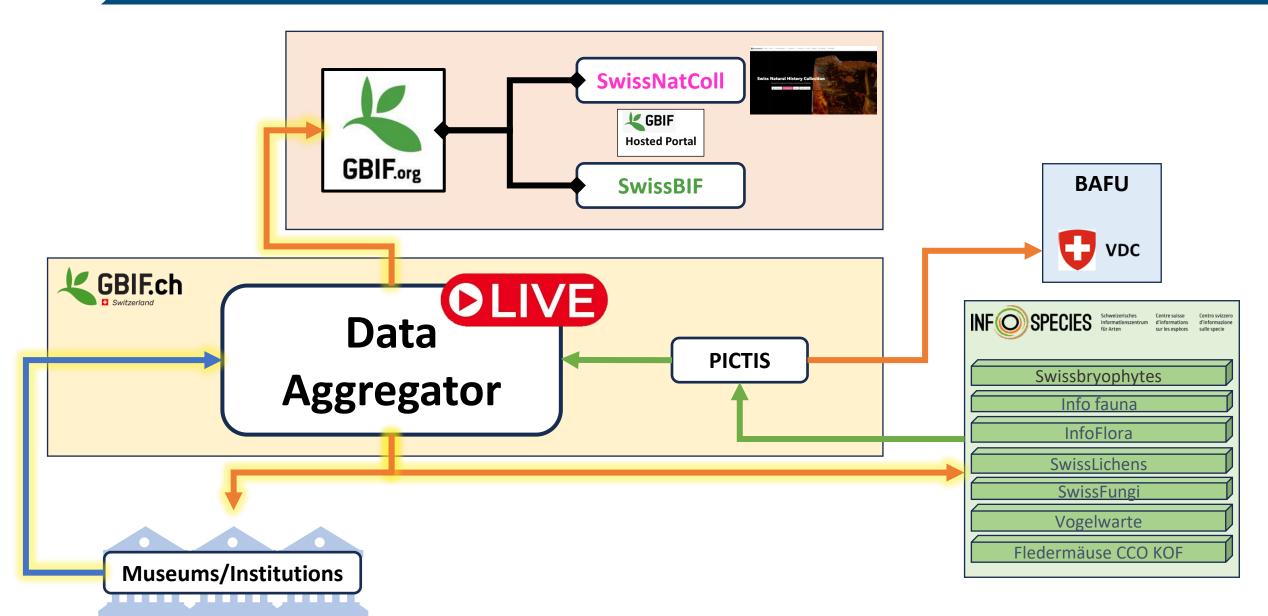
Introduction

Log In

Data preparation

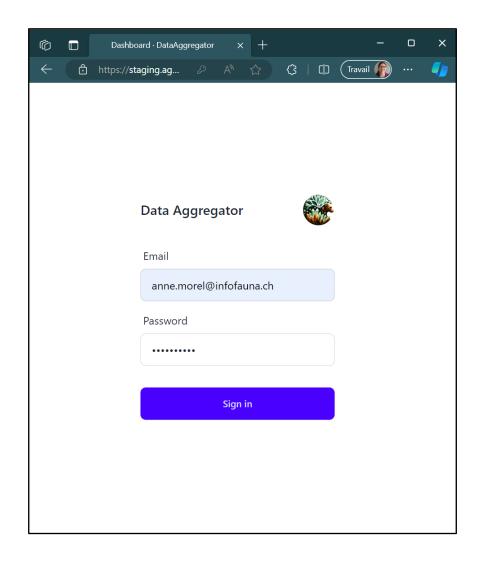
ZOOM sessions







Access to the Data Aggregator



Current URL link to the Data Aggregator

https://staging.aggregator.scnat.zebbra.ch/sign-in

To add the first user for your institution

contact-swissnatcoll@infofauna.ch

- Full name
- Email
- Institution



Data Administrator & Collection Digitizer

For all users and roles, the following rules apply:

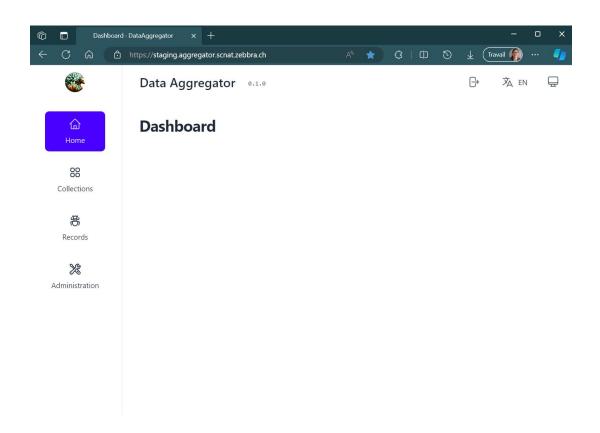
- One institution can have more than one user / role
- One **user** is attributed to **one and only one institution**
- One user can have more than one role
- A user profile is specific to an individual and must not be shared with others

Tasks	Data Administrator	Collection Digitizer	Admin
Users -add/edit/remove Collection Digitizer	×	×	✓ all
Users -add/edit/remove Data Administrator	X	own institution	
Collections -add/edit	×	own institution	
Collections -view	own institution	own institution	
Dataset -upload/mapping/encode/export/send for validation/publish		×	



! Page of one and only one institution! Home (planned)

- Link to the Step-by-step guide page
 (https://swissnatcoll.hp.gbif-staging.org/en/how-to-publish-data)
- Other elements TBD

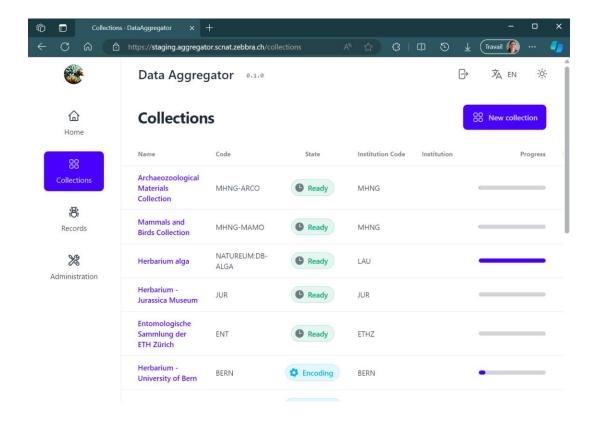




! Page of one and only one institution!
Home

Collections

- List of all collections already added on the Aggregator
- State and general overview
- Last update timestamp
- Button to add more collections (source: GRSciColl (gbif.org))





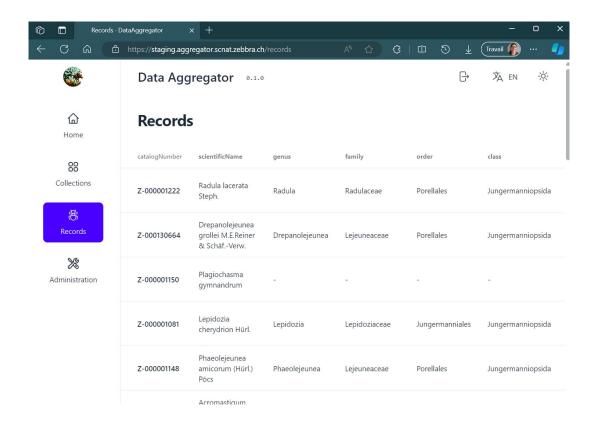
! Page of one and only one institution!

Home

Collections

Records

- List of all imported records for all collections
- State and general overview
- Last update timestamp





! Page of one and only one institution!

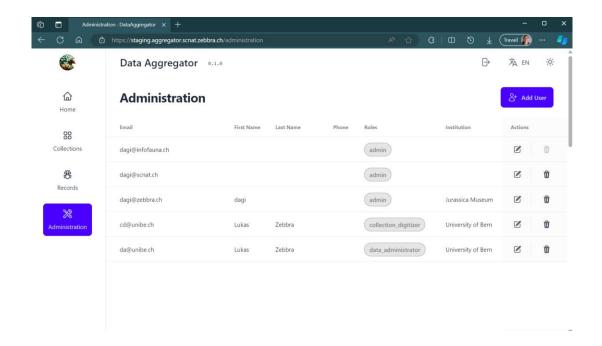
Home

Collections

Records

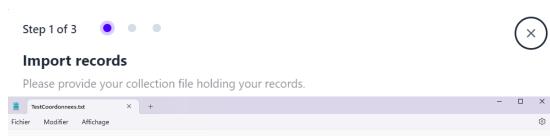
<u>Administration</u> (only for Collection Digitizer)

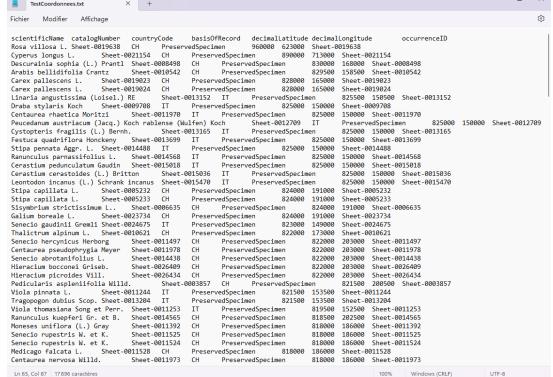
- List of all users of your institution
- Button to add more Data Administrator users





Upload file characteristics





Mandatory

- UTF-8 encoded file
- .csv, .tab, or .txt file
- 1 line = 1 record
- Unique catalog number
- Formatted fields

Facilitating

- Darwin Core headers
- Prioritization
- Tab separation



Fields mandatory and recommended

Mandatory

- catalogNumber
- scientificName

+ minimal recommended (MIDS 0-1-2)

- institutionCode
- partOfOrganism*
- preparations
- yearCollectionEntrance*
- collectionCode
- collectionID
- eventDate (OR day, month, year, endOfPeriodday/m/y*)
- recordedBy
- typeStatus
- locality
- decimalLatitude**
- decimalLongitude**

+ minimal expected (MIDS 2-3)

- verbatimElevation/verbatimDepth
- continent
- higherGeography
- country
- stateProvince
- county
- associatedMedia
- recordNumber
- basisOfRecord
- geodeticDatum
- coordinateUncertaintyInMeters
- swiss coordinates x (LV95)*
- swiss coordinates y (LV95)*

^{*}borrowed terms (MIDS, GBIFch), **only for WGS84 coordinates



Useful links

- Step-by-step tutorial: https://swissnatcoll.hp.gbif-staging.org/en/how-to-publish-data
- Data Aggregator overview : https://swissnatcoll.hp.gbif-staging.org/en/data-aggregator-about/
- Data Aggregator FAQs: https://swissnatcoll.hp.gbif-staging.org/en/data-aggregator-faqs (is planned to be enriched by your questions too)
- Darwin Core summary: https://swissnatcoll.hp.gbif-staging.org/en/data-aggregator-faqs#what-is-darwin-core
- Examples with real museum specimens (more coming soon)
 - Mollusca: https://swissnatcoll.hp.gbif-staging.org/en/label-mollusca



Take-Home Message

T Start small

- Check the mandatory fields in your database to make sure they are standardised
- Then move on to the next level of minimum fields
- Test first with a small file and then proceed with a bigger amount of records
- Error messages are a good point
 - During upload, error messages show inconsistencies in your database
 - They are opportunities to correct them
- Expected changes in the interface and functionalities
 - Development is still ongoing, bugs are currently being solved
 - Your contributions are essential for this user-oriented tool