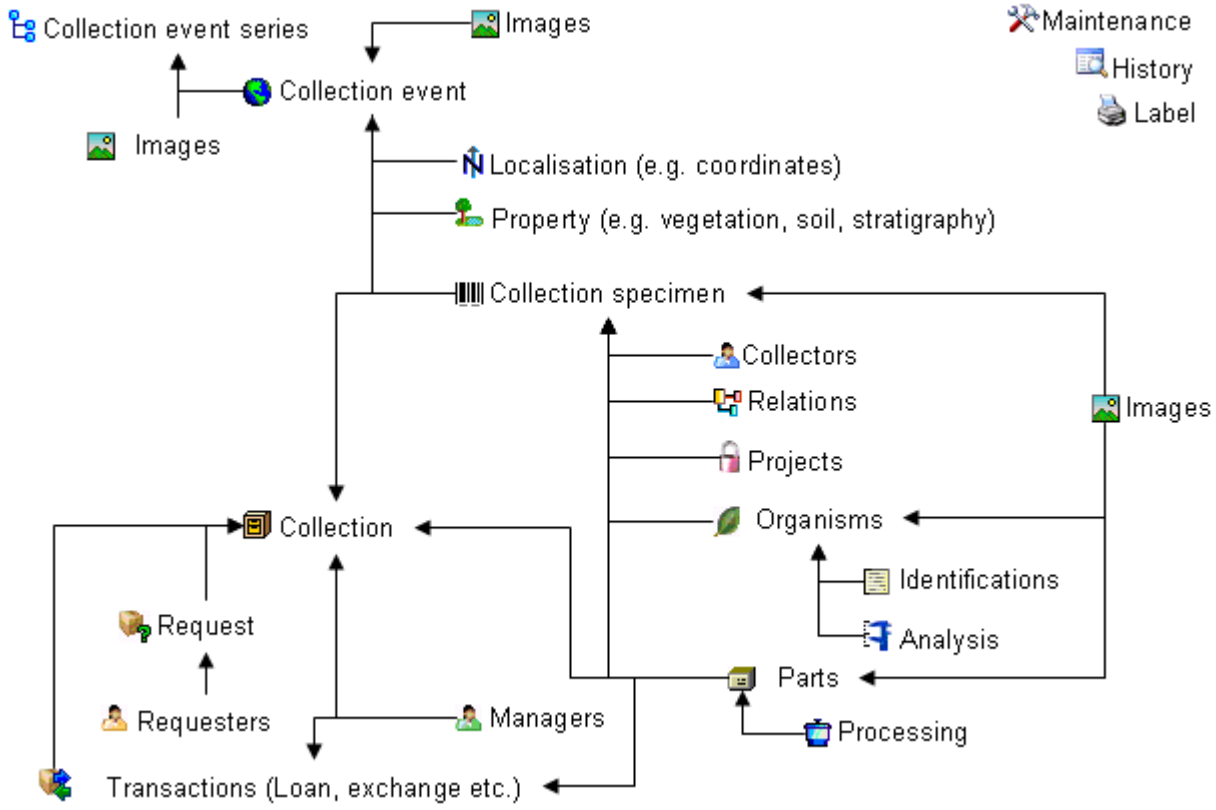
DiversityCollection

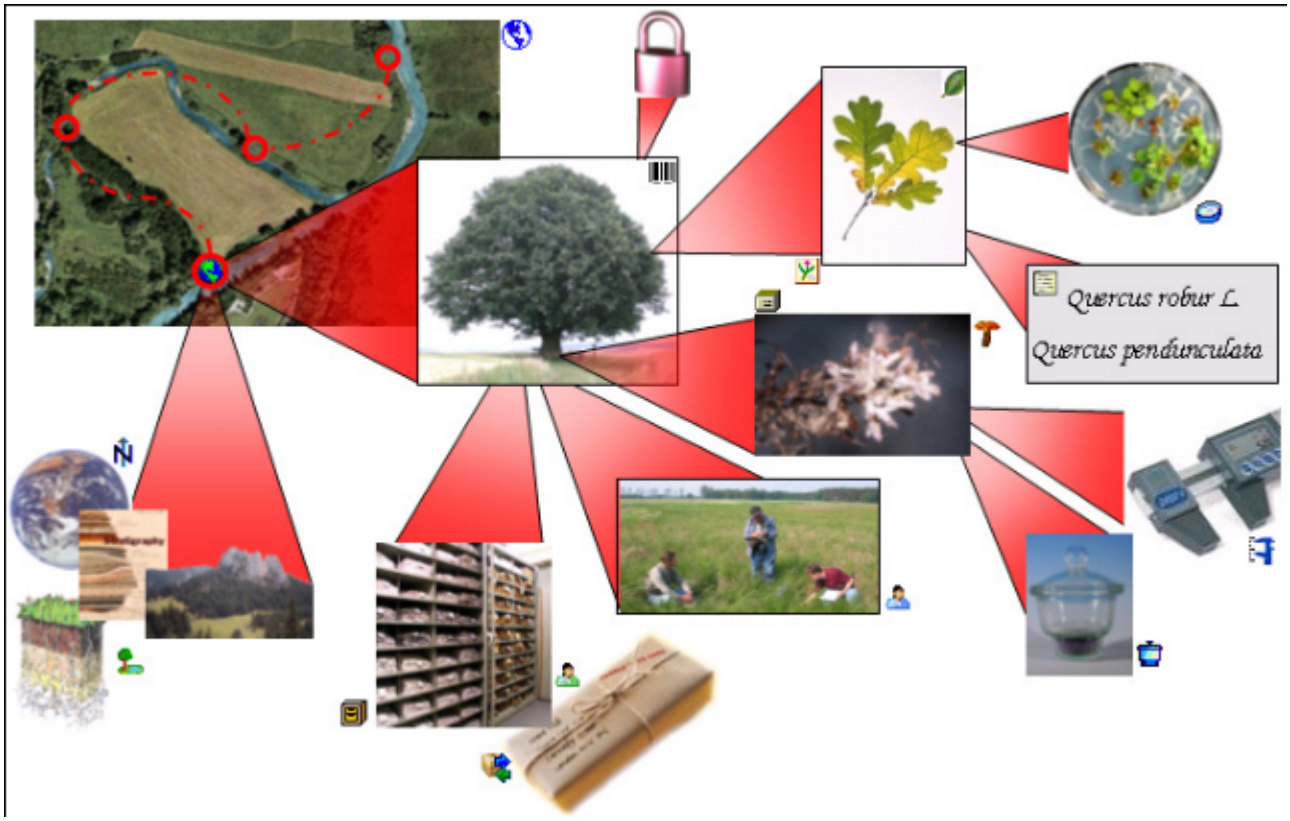
DiversityCollection (Version 2) is part of the database framework Diversity Workbench. Within this framework the application DiversityCollection is confined to the management of specimens in scientific collections. In this context it is designed to documente any action concerning the collection, storage, exchange and treatment of specimens in a collection and is also appropriate to store observation data. DiversityCollection is distinguished from other collection management systems by its focus on biological relations between organisms collected together as one or more specimens (e.g. host, parasite and hyperparasite, symbionts etc.). Any module within the Diversity Workbench is focused on a specific data domain. DiversityCollection keeps only data connected with the handling of collection specimens and observations. Data of other realms like e.g. taxonomy are handled in separate modules. For an overview of the available modules see Diversity Workbench. DiversityCollection might also be used as a stand-alone application.

The image below gives you an overview of the main parts, relations and functions of DiversityCollection



The image below gives an overview for some typical data depicted in DiversityCollection together with the symbols used throughout the program. A typical specimen may have been collected at a collection event during an expedition. The site of the collection event may be localized and characterized. The collectors collected twigs of the plant and fungi from the roots. They store the samples as herbarium sheets and specimens preserved by other methods in a collection. The manager sent some of the samples for which he had a request from a requester to another collection. The samples were collected as part of a project. Certain parts were cultivated, analysed and processed. The organisms on specimen were identified several times. Images were taken for the event series, the collection event, the collection specimen as well as organisms

and part of this specimen.



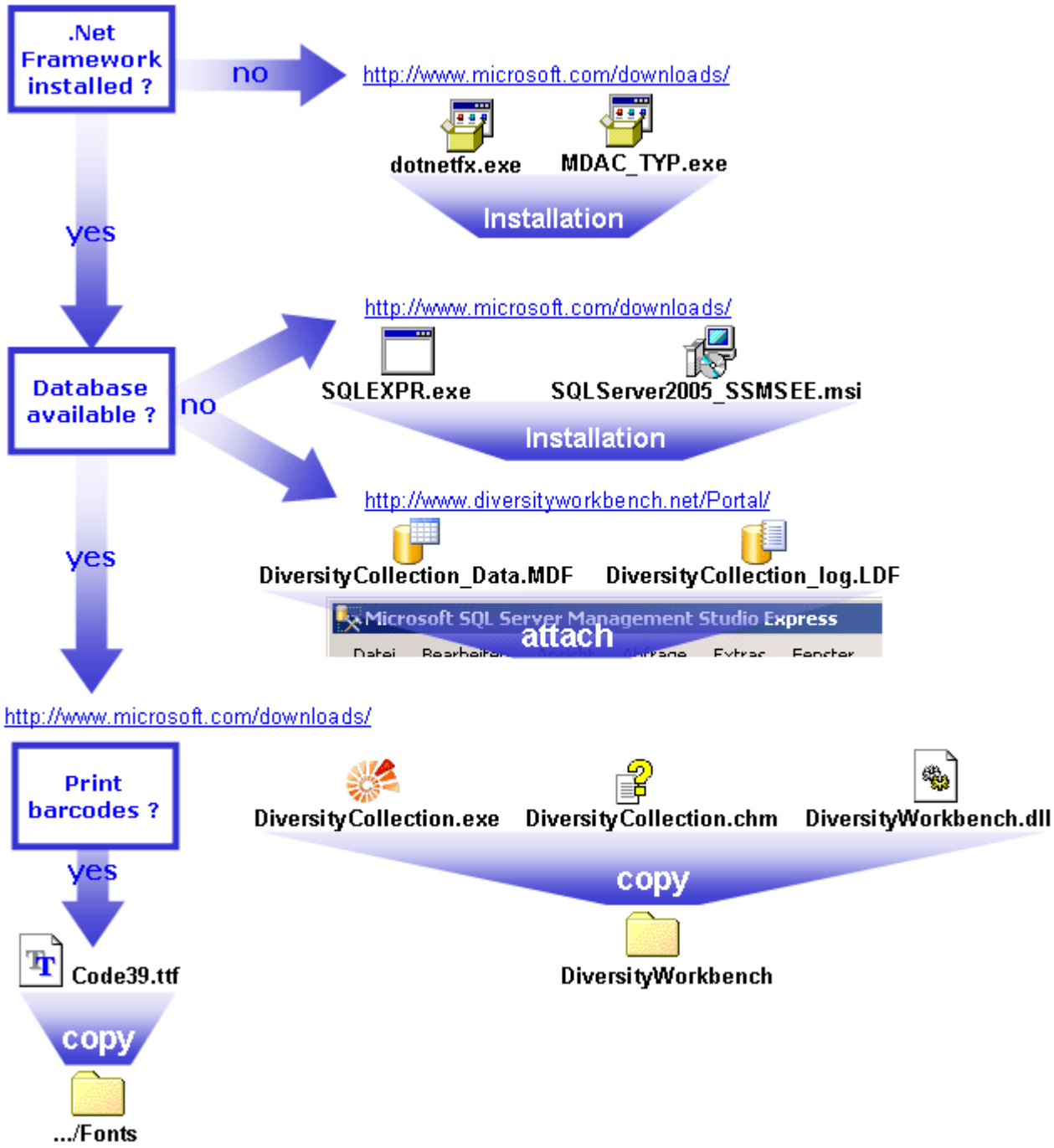
DiversityCollection 2 is based on Microsoft SQL-Server 2005 and the .Net Framework, Version 2.0.

For licence and copyright see the licence section.

Installation

To run DiversityCollection, you need the database and the client. All parts are free and can be downloaded from http://www.microsoft.com/downloads/ and http://www.diversityworkbench.net/Portal/.

The image below gives an overview of the installations and files needed.



Client

The client is based on the .Net framework version 2.0 from Microsoft. If not already present, you have to install the framework first. Download and install the Microsoft .NET Framework (e.g. dotnetfx35.exe - start the program and follow the installation instructions (see

http://www.microsoft.com/downloads/ for the latest versions). Version 2.0 is compatible with older operating systems like Windows 2000. For later operating systems use the latest version available.

Download the files for DiversityCollection from http://www.diversityworkbench.net/Portal/ provided as a zip archive. Copy all files (DiversityCollection.exe , DiversityCollection.chm , DiversityWorkbench.dll) into your DiversityWorkbench directory. To print barcodes of the font Code 39 the file Code39.ttf must be copied to the folder C:\WINNT\Fonts.

After the installation make shure to get the latest updates from http://windowsupdate.microsoft.com/.

Database

For the installation of a local database see the section Installation of the database.

Menu

Overview of the menu in DiversityCollection

Connection

Database ...	Choose one of the databases available on the server. Only those databases will be listed to which the user has access permission
Module connections ...	Edit the connections to the other modules within the Diversity Workbench.
Transfer previous settings	Transfer the settings of a previous version
Quit	Quit the application and stop all processes started by the application

Query

Show query	Show or hide the query list
Predefined queries	Under this menu entry all predefined queries will be listed
Scan mode	Change to the scan mode to open the dataset of a specimen by scanning the barcode
Grid mode	Change to the grid mode to edit the dataset in a list

Data

Import	
Specimen scans ...	Import scans of specimen labels
Import list ...	Import tab-separated lists
Export	
Export list ...	Export a tabulator separated file with the data of the specimen
XML ...	Export data as a XML-file according to ABCD

Administration

Analysis ...	Administration of the analysis methods used for the specimens
Collections ...	Administration used for the collections
Customize display ...	Customizing the display of the window, e.g. the material categories and taxoniomc groups that should be visible when creating a new entries
Maintenance ...	Maintenance of database entries especially if connected to other modules
Processing ...	Administration of the processing procedures applied in the collection
Queries ...	Creating and editing predefined queries
Transaction management	Management transactions, managers, loans etc.
Transactions ...	Administration of the transactions, e.g. loans, exchange etc.
Expired loans ...	Administration of expired loans. This menu entry will appear when there are expired loans in collections where the current user is a curator
Loan requests ...	Administration of loan requests for the collections a user is an collection manager. This menu entry will appear when there are loan requests for the managed collections of the current
My requests ...	Administration of the loan requests of a user. This menu entry will appear when a user placed requests for specimen
Requesters ...	User having the right to place requests for specimen of a collection

Managers ...	Administration of the users that manage collections and are e.g. responsible for the administration of the transactions
User ...	Administration of the users and their permissions in the database
Versions ...	Administration of the versions of client and database

Help

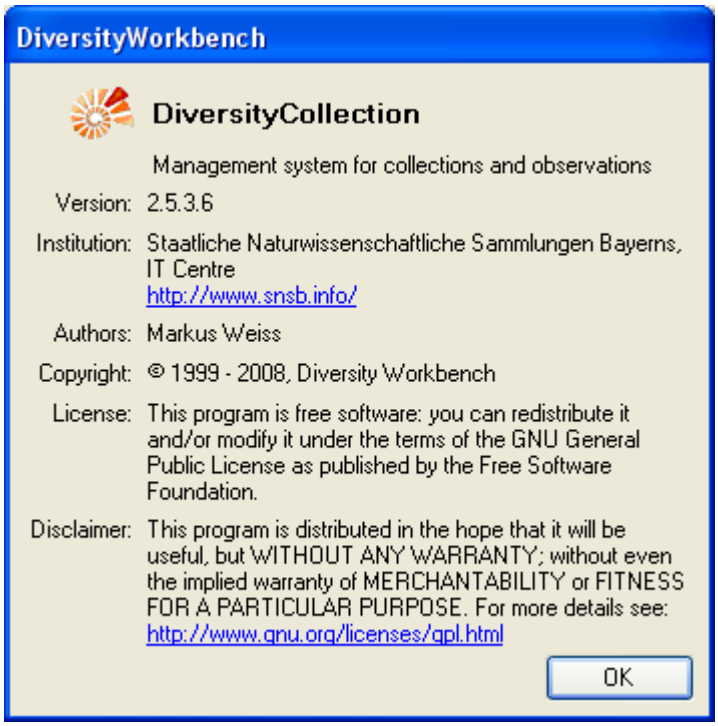
Manual	Opens the online manual
Feedback ...	Opens a window for sending feedback
Info	Show the version of the program and corresponding information

Manual

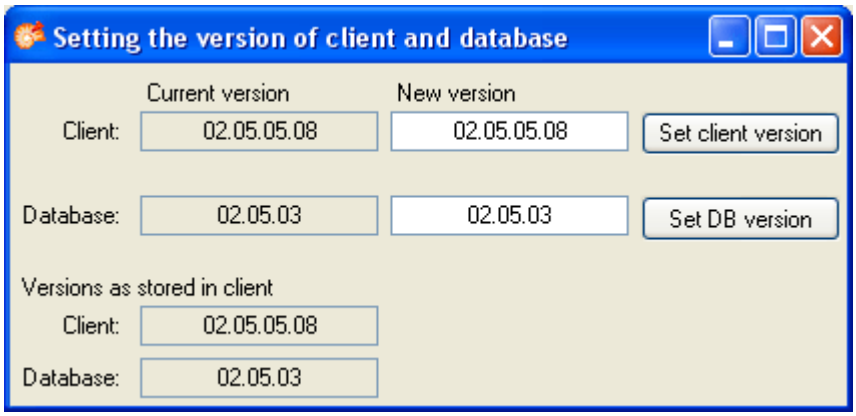
The online manual DiversityCollection.chm must be placed in your application folder, together with the application DiversityCollection.exe and the library DiversityWorkbench.dll. To get information to any topic in the application DiversityCollection and open this manual, just click on the field you need information about and press F1. To open the manual from the menu, choose **Help ->** **Manual**.

Version

For information about the version of the client application choose **Help**, **Info...**

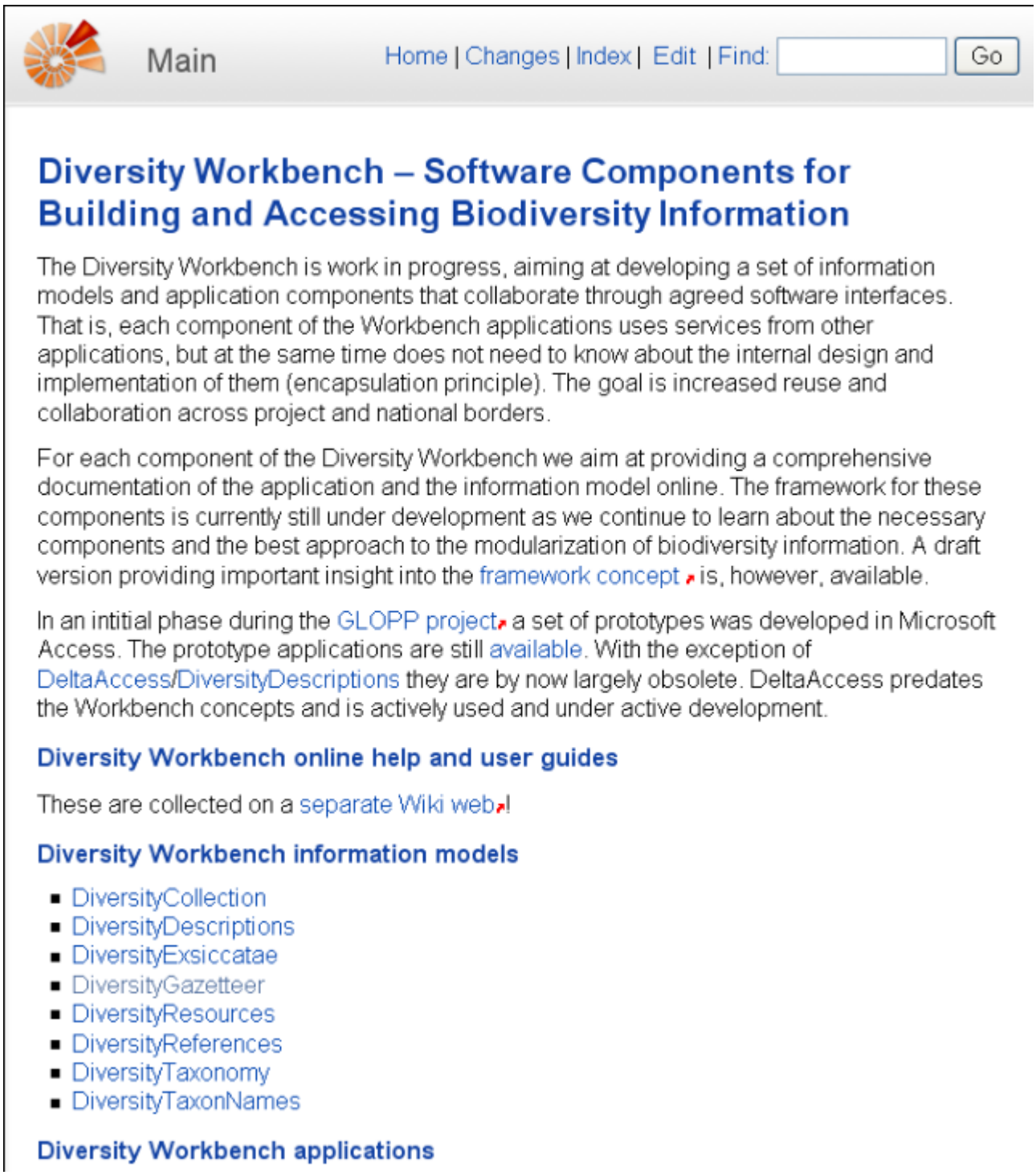


The current version in the example above is 2.5.3.6. As an administrator, you can set the versions of the database and the client. Choose **Administration** - **Versions** from the menu. A form as shown below will open, giving you an overview of the version settings.



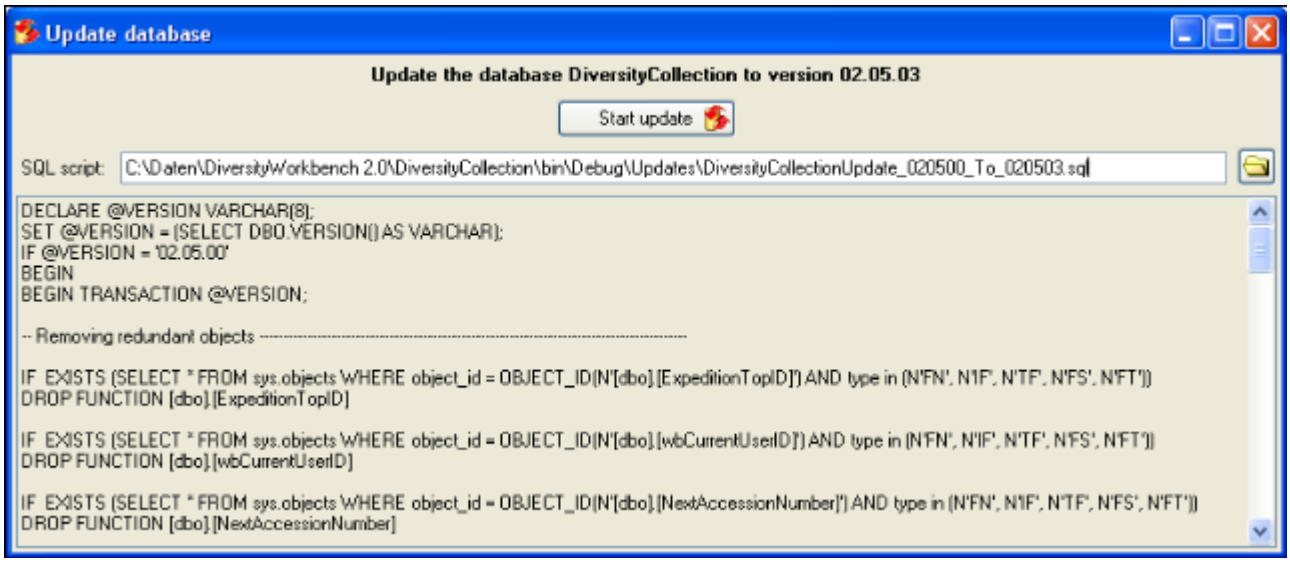
Update of database and client

DiversityCollection is still in development. Therefore regular updates for the database and the client will be provided. When you start the program and connect to a database, the program will check if it is compatible with the database or if the database needs an update. In any of these cases an **update** entry in the menu will appear. If a new version of the client is available this menu will contain an **update client ...** entry. Click on it to open the webpage where you can download the cient as shown below.



If you are the owner of the database (Database role = dbo) and the database needs to be updated, the menu will contain a **update database ...** entry. Select this entry to open a window as shown below to run the provided update scripts, delivered with the client software.

These scripts needs to run consecutively, so e.g. to update from version 2.5.1 to 2.5.4 you either have to run the script DiversityCollectionUpdate_020501_To_020504 or the scripts DiversityCollectionUpdate_020501_To_020502, DiversityCollectionUpdate_020502_To_020503 and DiversityCollectionUpdate_020503_To_020504. The program will guide you through these steps and check for the scripts. All you have to do is click the **Start update** button.



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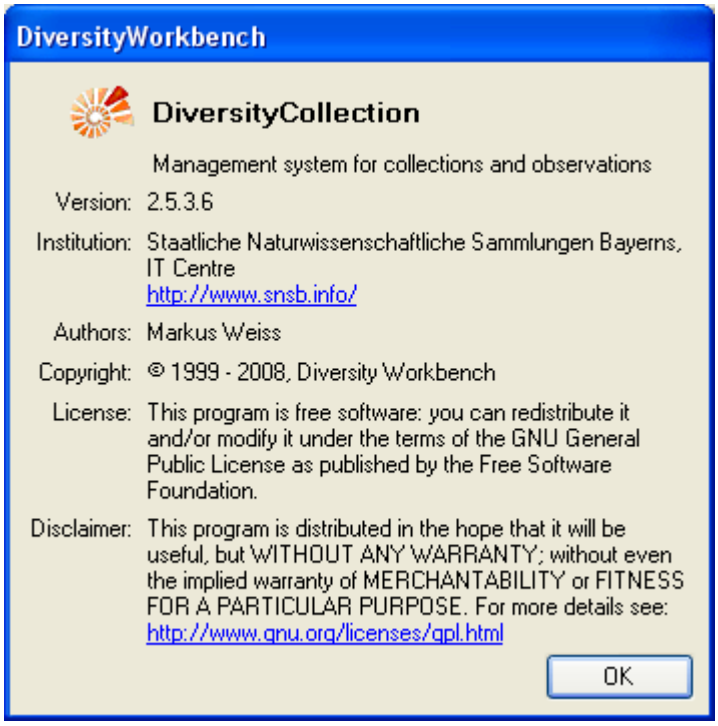
For information about the license of the client software choose **Help**, **Info...**



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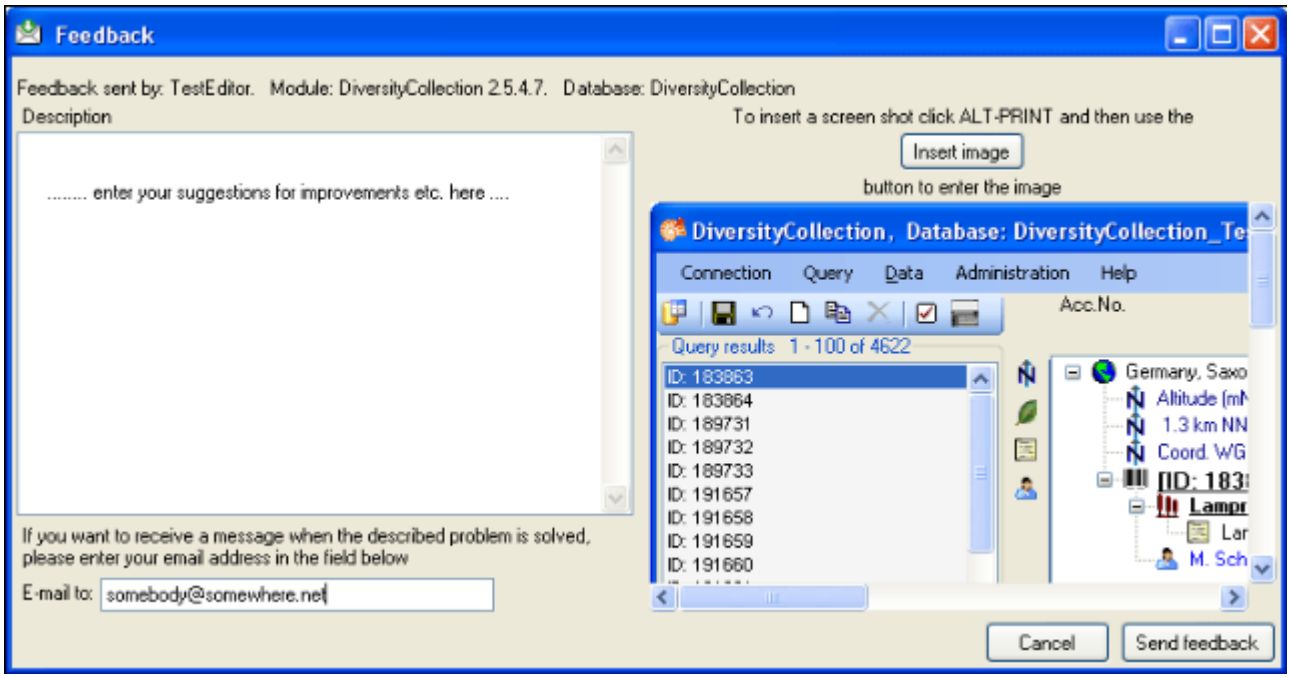
The client software is distributed in the hope that it will be useful, but WITHOUT ANY WARRANTY; without even the implied warranty of MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the GNU General Public License (GPL) for more details.

Feedback

If you have suggestions for improvement, need any changes in the program or encounter an error you can give a feedback to the administrator. Click on the [**ALT**] and [**PRINT**] buttons to get a screen shot of your current form. After creating the screen shot choose **Help - Feedback** from the menu to open the feedback window as shown below.

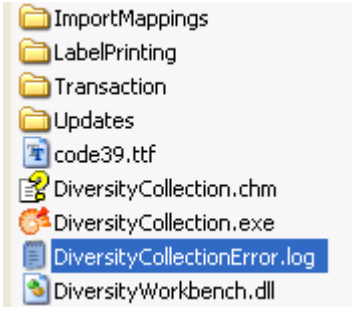


Click on the Insert image button to insert the screen shot and give a comment about your problem. Then click on the Send feedback button to send your feedback to the administrator. If you want to receive a message, when the problem you described is solved, please enter you e-mail address in the field under the descrption.

In case you do not have access to the central database for the feedbacks, the program will open your mail client to send an e-mail. In case of bugs in the program it would help if you attach the file **DiversityCollectionError.log** located in your application directory (see image below).

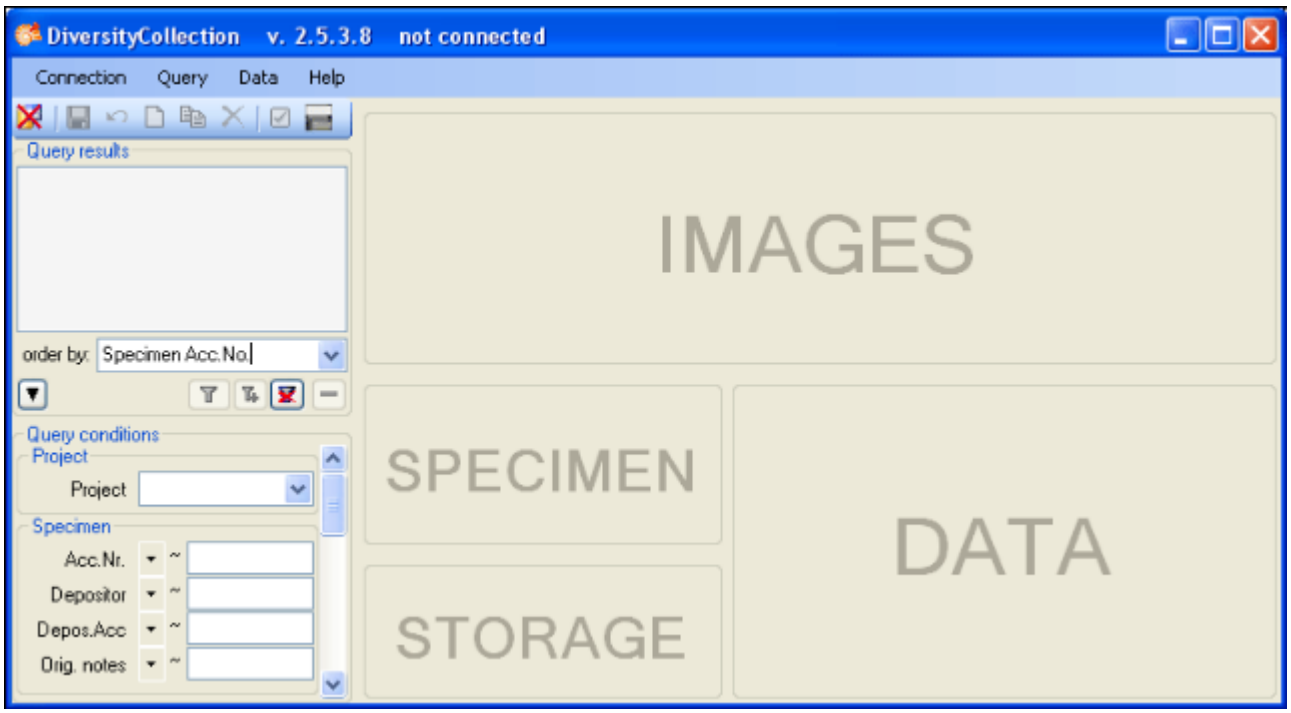
Error logging

If any error messages show up through working with the application you can find further details concerning the part of the application where the error occured and the parameters involved in the file **DiversityCollectionError.log** located in your application directory.

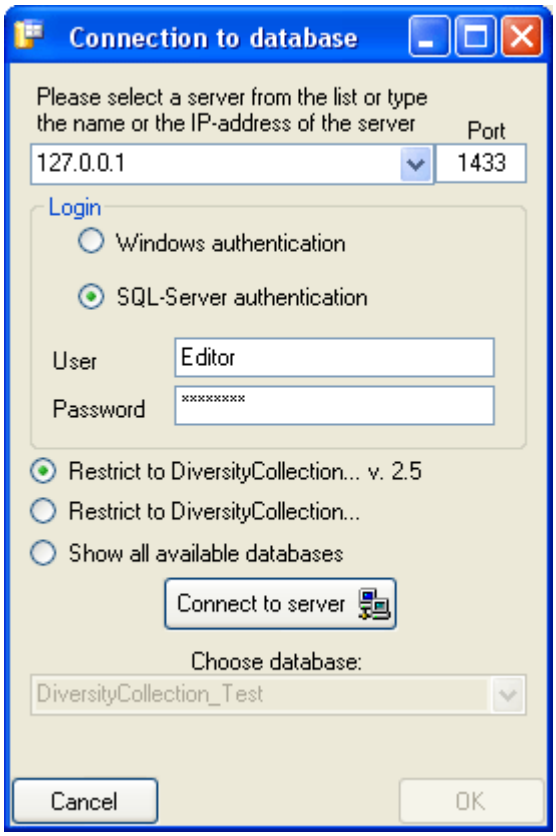


Tutorial - first steps

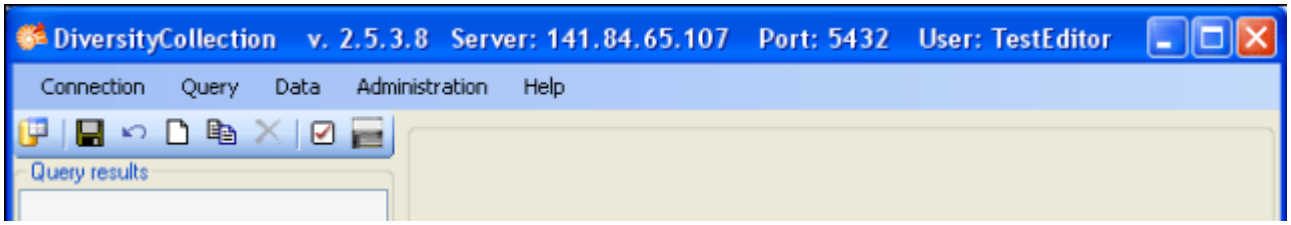
This tutorial will guide you through the first basic steps to enter a single dataset in DiversityCollection. After the installation, make sure, you have access to the database. To start the programm, double click on the DiversityCollection.exe in the directory where you copied the files of DiversityCollection. The main window will open.



If you open this window for the first time, you have to connect to the database. Click on the button or choose **Connection -> Database...** from the menu. A window will open where you can enter your account information and choose the database (see image below, for further informations see database access).



After connecting to the server and choosing a database click on the **OK** button to return to the main form. As indicated by the symbol in the right upper corner, you are now connected to the database. In addition the header of the window shows your current login informations (see below).



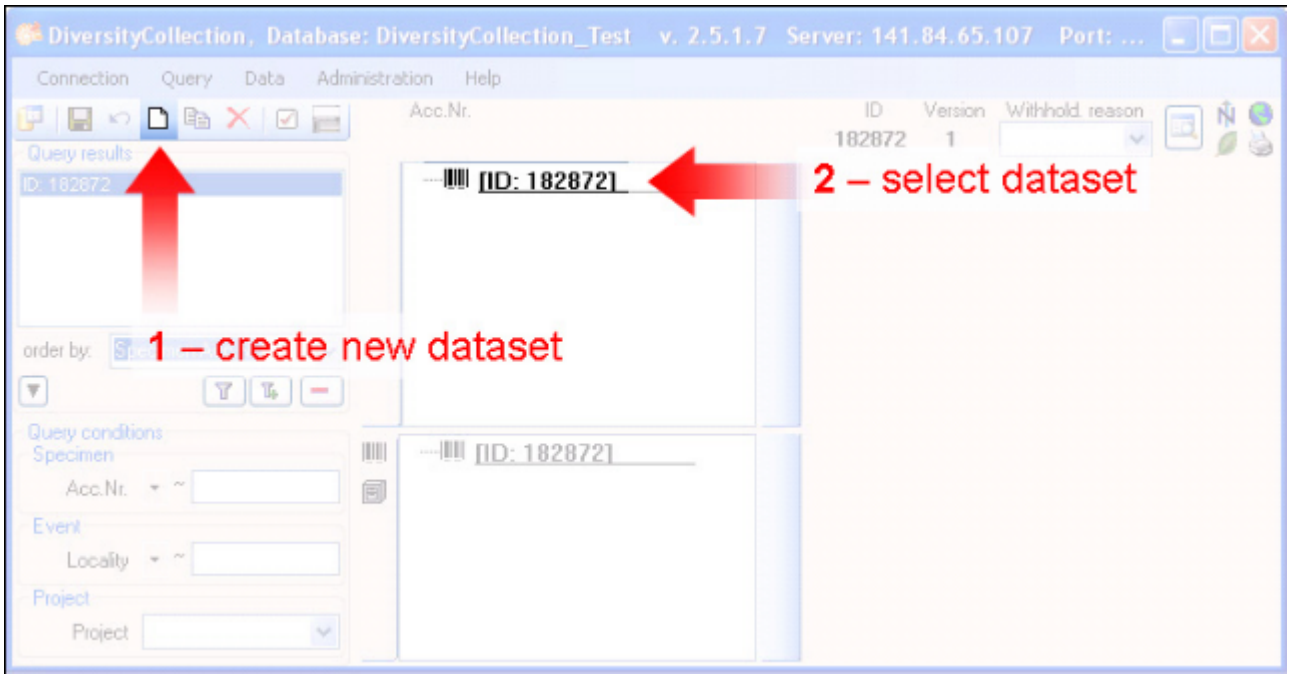
This tutorial is continued in the sections listed below.

- section Collection event
- section Collection specimen
- section Collection specimen relations
- section Organisms and identifications
- section Specimen parts and storage

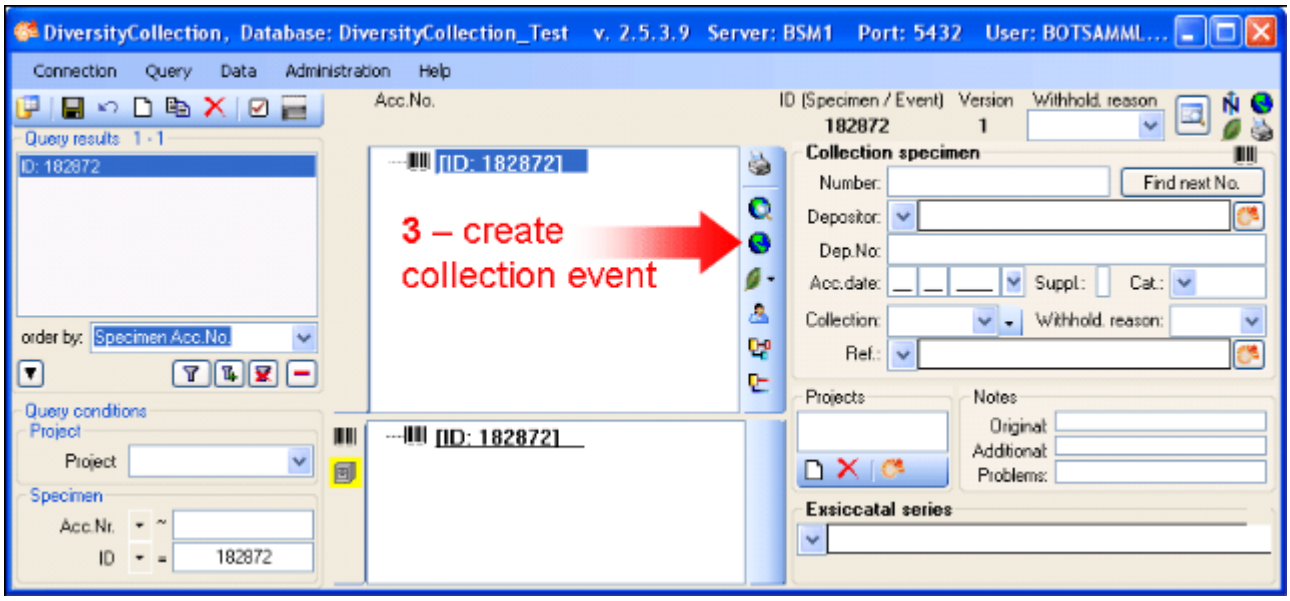
Tutorial - collection event

To enter a new dataset, click on the button in the upper left panel (see point **1** in image below). In case there are accession numbers in the database, the software will ask whether you want to take the next free number (see accession for further information). If you click on the OK button, the program will try to find the next free accession number starting with your current accession number.

Finally you will find a new entry in the specimenlist on the left and the trees for the specimen (see point **2** in image below) and the storage as shown below.



If you did not specify an accession number the entries will look like in the image above, otherwise the accession number will be shown. To enter your data for the specimen, click on the entry in the upper tree as shown below. This will open the fields where you can enter the details for the specimen in the area right from the tree.

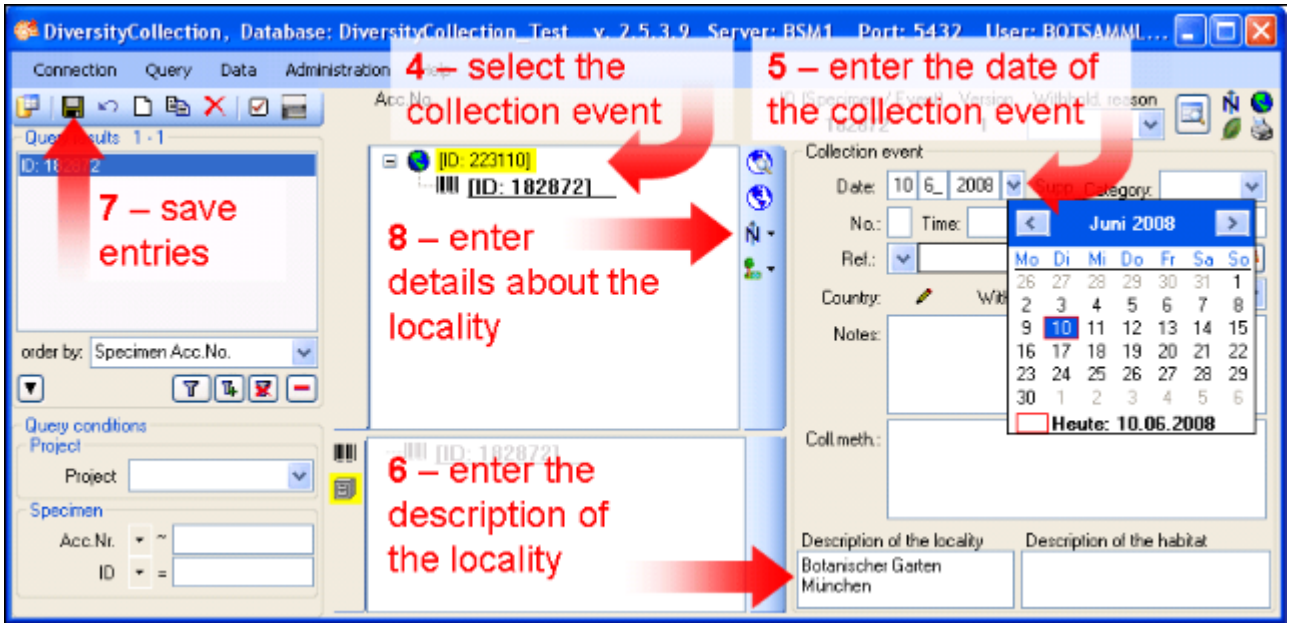


In the panel in the middle of the form, several buttons will appear where you can enter additional informations to this specimen. To enter information about the collection event (when and where the specimen was collected) click on the button (see point **3** in image above). This will add an entry for the collection event in the upper tree as shown below. Select this entry in the tree (see point **4** in image below), to open the fields for the collection event. To see the projected contents of any data field, simply place your mouse in the field. A explanation will appear like for the field **Description of the locality**:

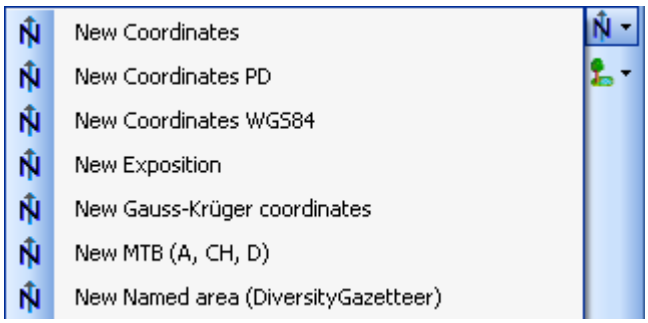
Locality description of the locality, exactly as written on the original label (i.e. without corrections during data entry)

See the event part for further details.

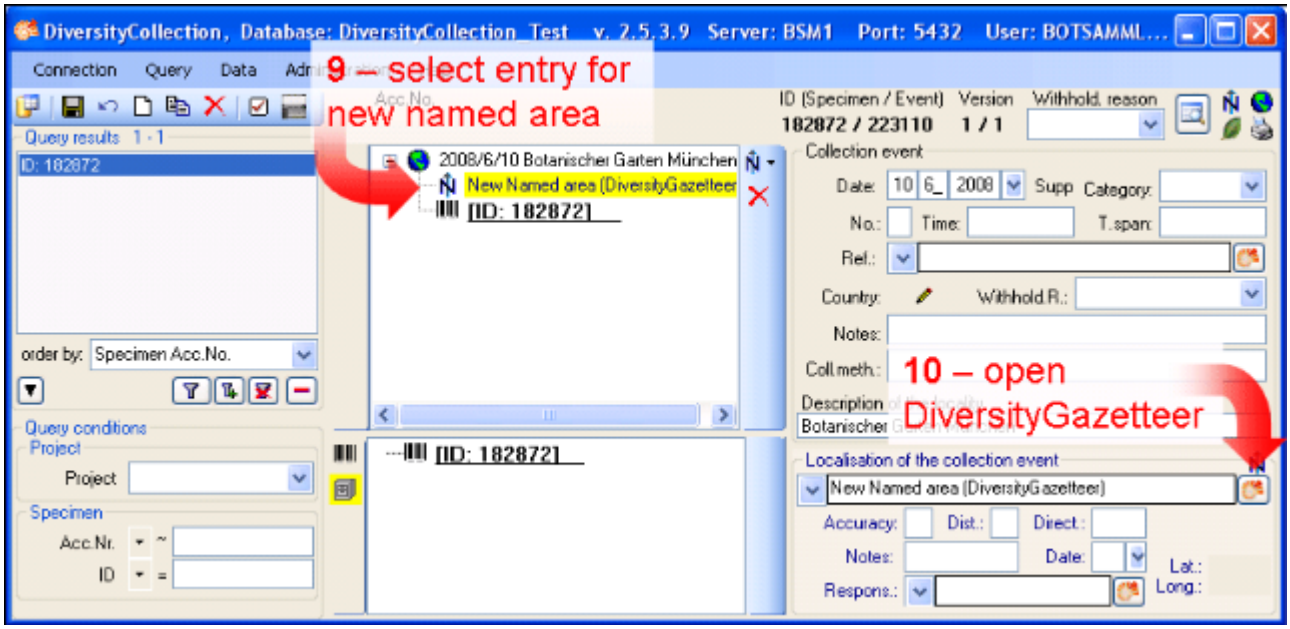
Then enter the date (see point **5** in image below) of the collection event. If you click on the drop down button as shown in the image below, a calendar will open where you can select the date. Then enter the description of the locality (see point **6** in image below). To store the data entered so far, click on the button (see point **7** in image below).



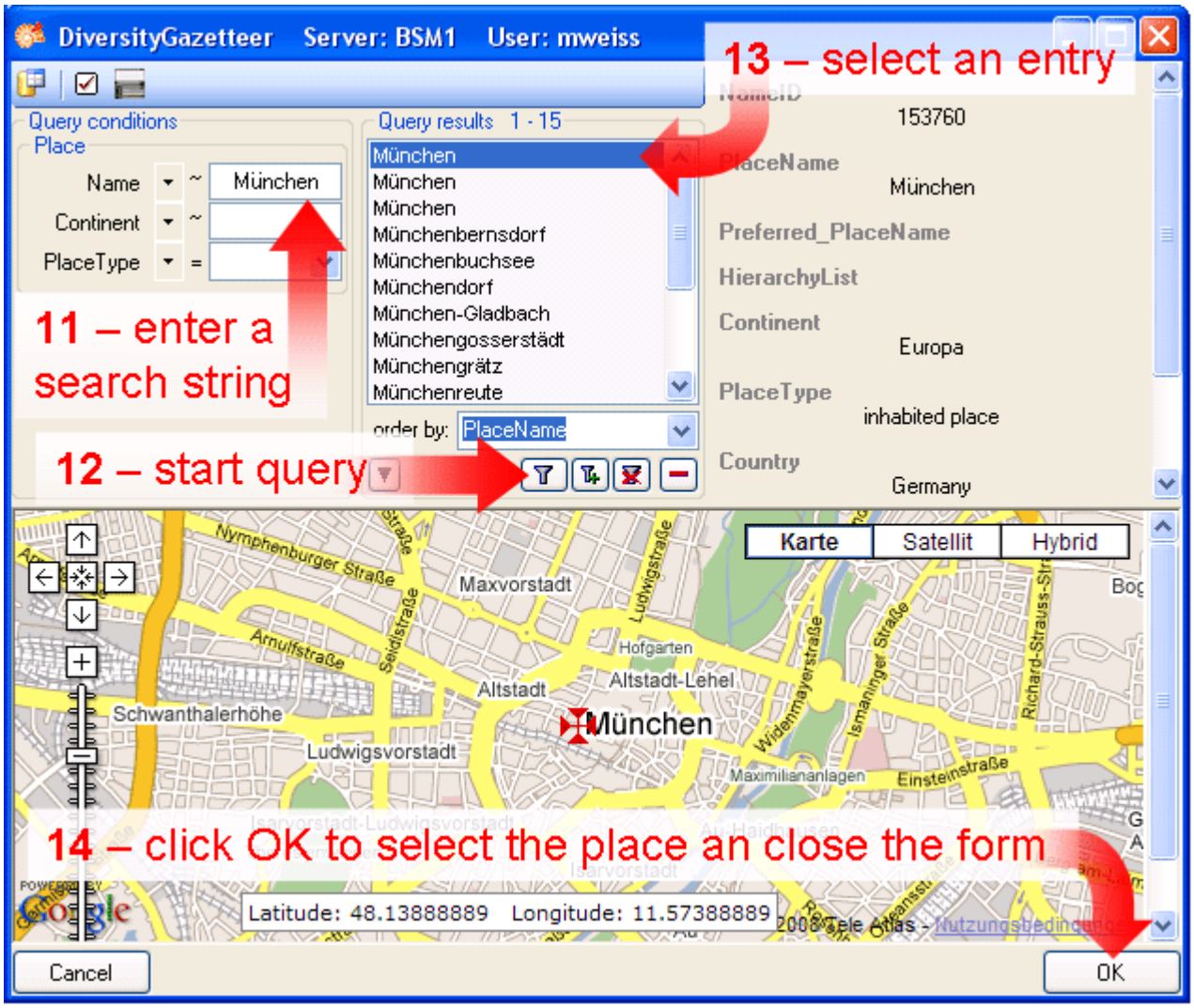
To enter coordinates, named places etc. click on the button (see point **8** in image above) and select an option from the list.



To enter a name of the place using the DiversityGazetteer, choose **New Named area (DiversityGazetteer)** from the list. In the overview tree in the middle of the window an entry will be inserted as shown below. Select it to open the detail fields for this entry (see point **9** in image below).



To open the connection to the DiversityGazetteer, click on the button (see point **10** in image above). A window as shown below will open. Enter a search string for e.g. the city you want to find (see point **11** in image below) and start the query with a click on the button (see point **12** in image below).

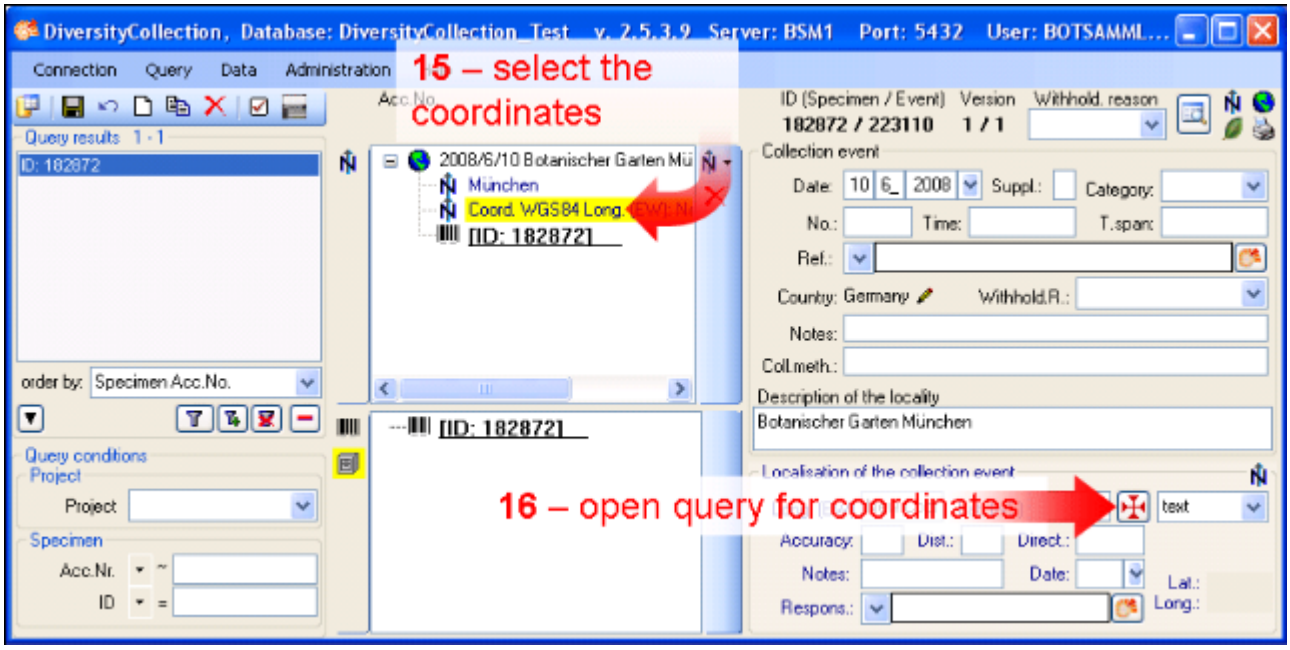


In the middle of the form, the results of the query will be listed. Select on of these. To guide you to the correct entry, details to this place are listed in the area right from the list and the base will show a map corresponding to the coordinates connected to this entry. After selecting the correct entry (see point **13** in image above) click on the OK button to return to the main window (see point **14** in image above). As shown below the data retrieved from the gazetteer will be written in several areas. Next to the name of the place the DiversityGazetteer provides the coordinates and the country as shown below.

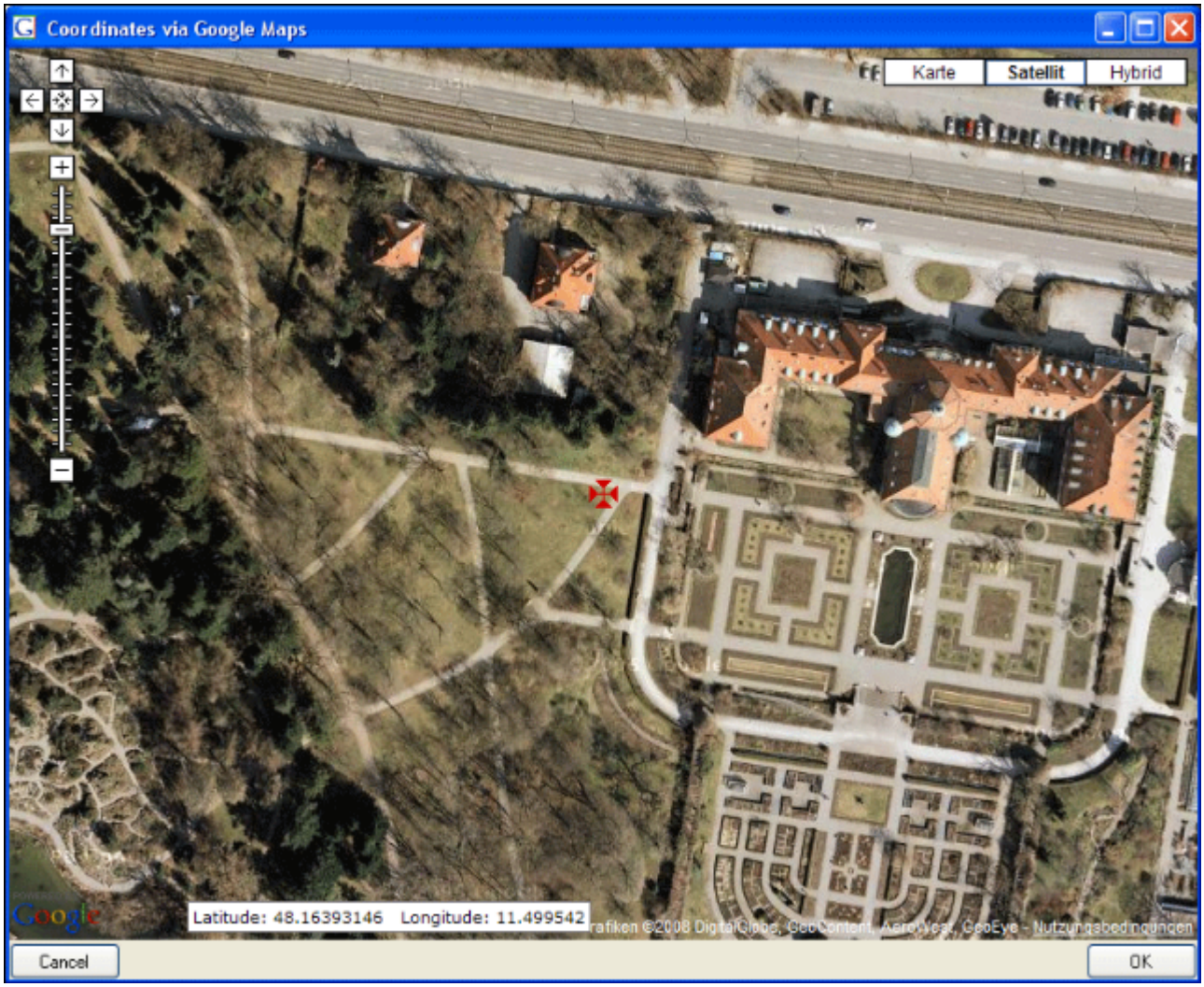
The area where you entered the name now changed to a locked state and will prevent you from changing the entry (see image below). Next to the field with the place you find the link to the external module. Double-click it for more details.

If you want to remove the connection to the external module click on the button. This will keep all entries (country, place, coordinates) but remove the connection to DiversityGazetteer.

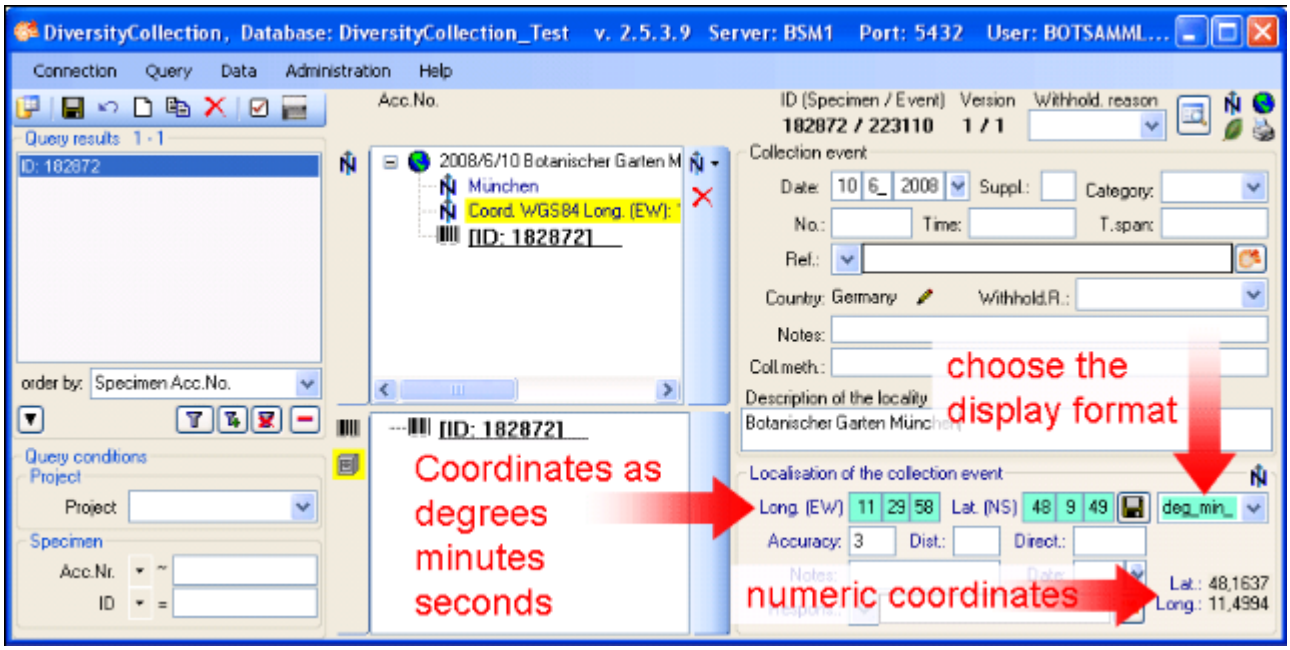
Now lets add the exact coordinates for the locality with the assistance of Google Maps. Click on the button (see point **8** above) and choose **New Coordinates WGS84** (Google Maps uses WGS84). This will add a new entry in the overview. Select it (see point **15** in the image below) to open the details for this entry. Here click on the button (see point **16** in image below).



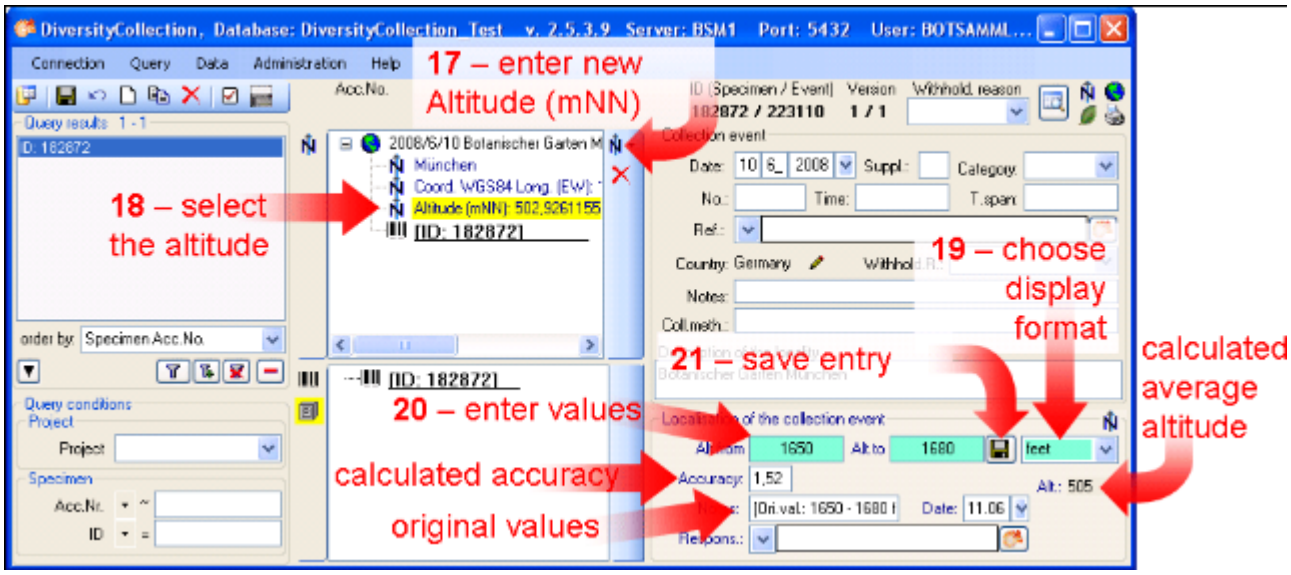
A window as shown below will open where you can set the coordintes simply by dragging the map with your mouse. The coordinates correspond to the center of the map, symbolized with the ✠. Click on the OK button to store the coordinates.



In the main window as shown below the coordinates will be stored at two positions - see image below. In the upper area, you can set the values an choose a different format for display as shown here the more familiar form with degrees, minutes and seconds. You change the values and click on the button, to change the original entry. In parallel the numeric values are stored in fields that can not be edited by the user (see below).



As a last information about the locality, we enter the altitude. Click on the button (see point **8** above) and choose **New Altitude (mNN)** (see point **17** in the image below). Then select the new entry in the overview to open the datafields (see point **18** in the image below). Lets suppose, you have only feet values available - so change the display format to feet (see point **19** in the image below). Enter your values (see point **20** in the image below) and click on the button (see point **21** in the image below) to save your entries. Now your values are converted to meter, the internal format of DiversityCollection. The program calculates an average value for the altitude and an accuracy in meter corresponding to the accuracy of your original values (see below). The original values of your entry are saved in the **Notes** field.

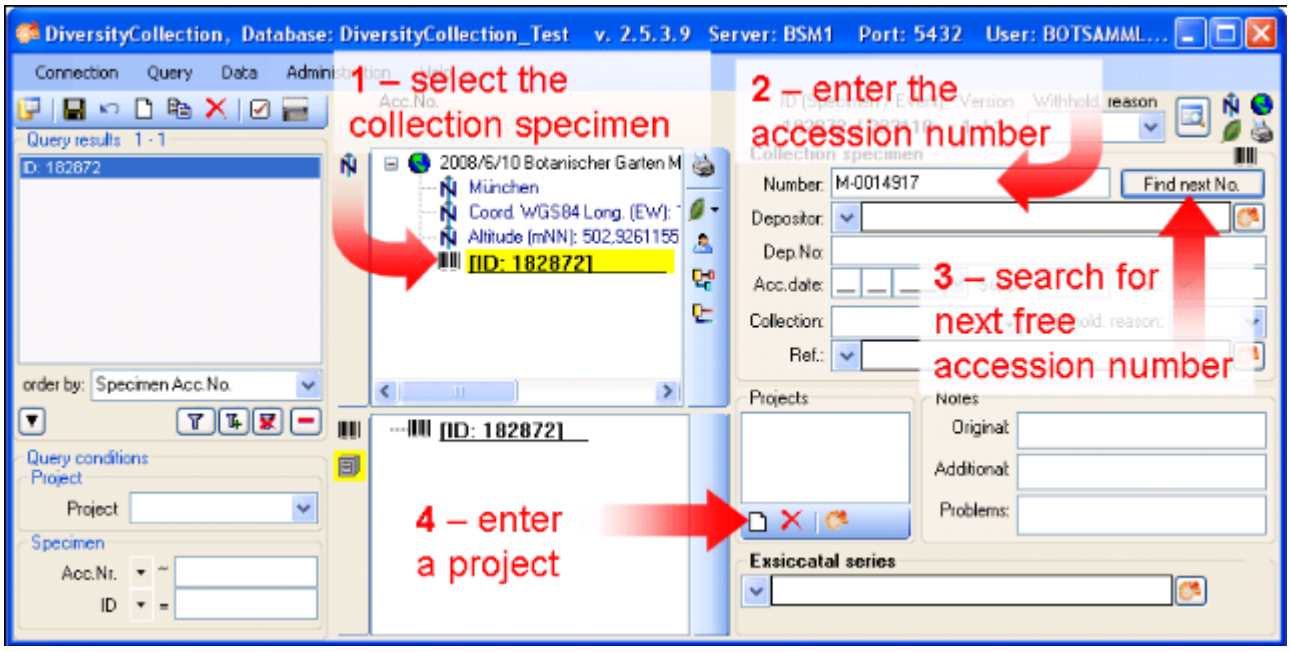


Next we turn to the collection specimen and set the accession number in the first section listed below.

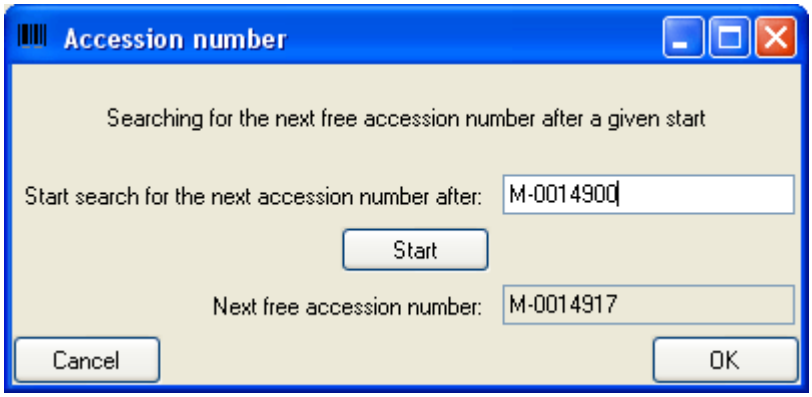
- section Collection specimen
- section Organisms and identifications
- section Specimen parts and storage

Tutorial - collection specimen

To set the accession number for the collection specimen, select it in the overview (see point **1** in image below) enter the accession number (see point **2** in image below).

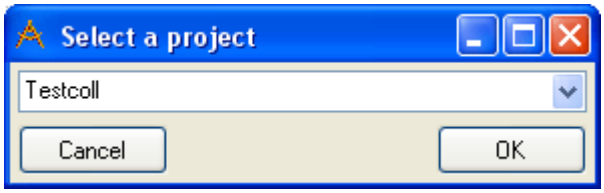


To search for the next free accession number, click on the corresponding button as shown above (see point **3** in image above). A window as shown below will open. Start the search for a free accession number. If the query ended successful, click **OK** to take the new accession number in your dataset.



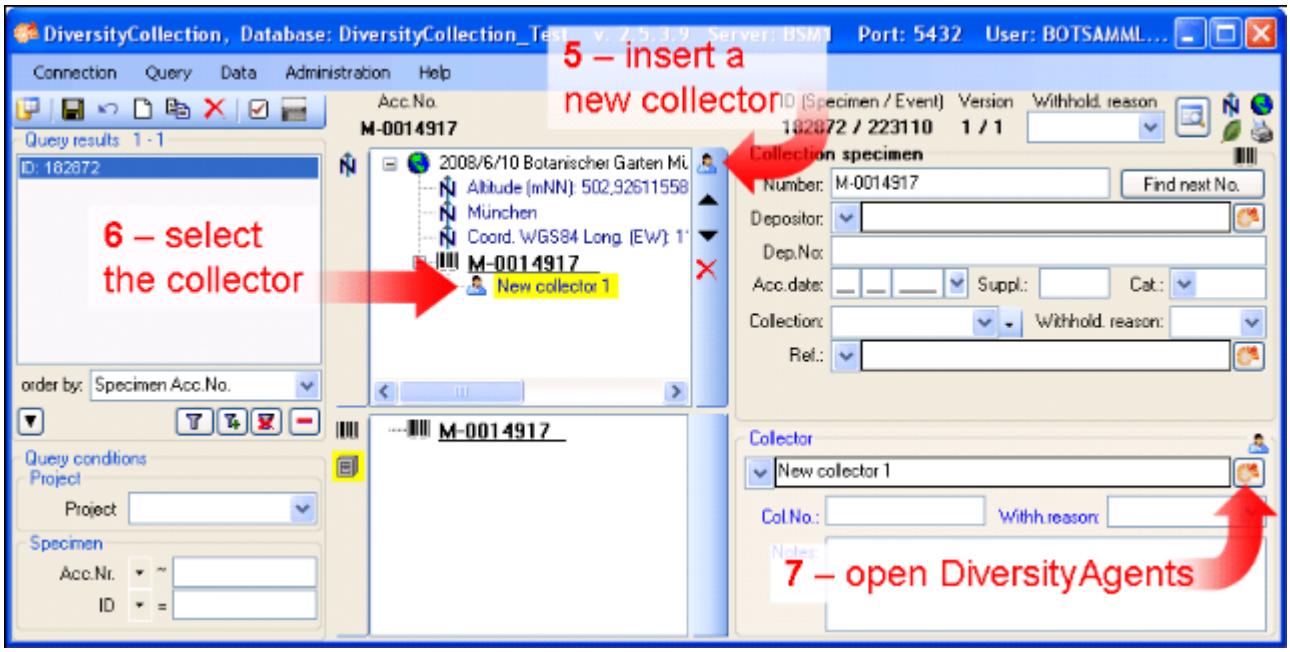
Project

To restrict the access to your dataset, add it to a project (click on the button - see point **4** in image above). A dialog will open where you can select a project from the projects defined in DiversityCollection (see image below). The project will then be added to the list of the projects for this specimen.

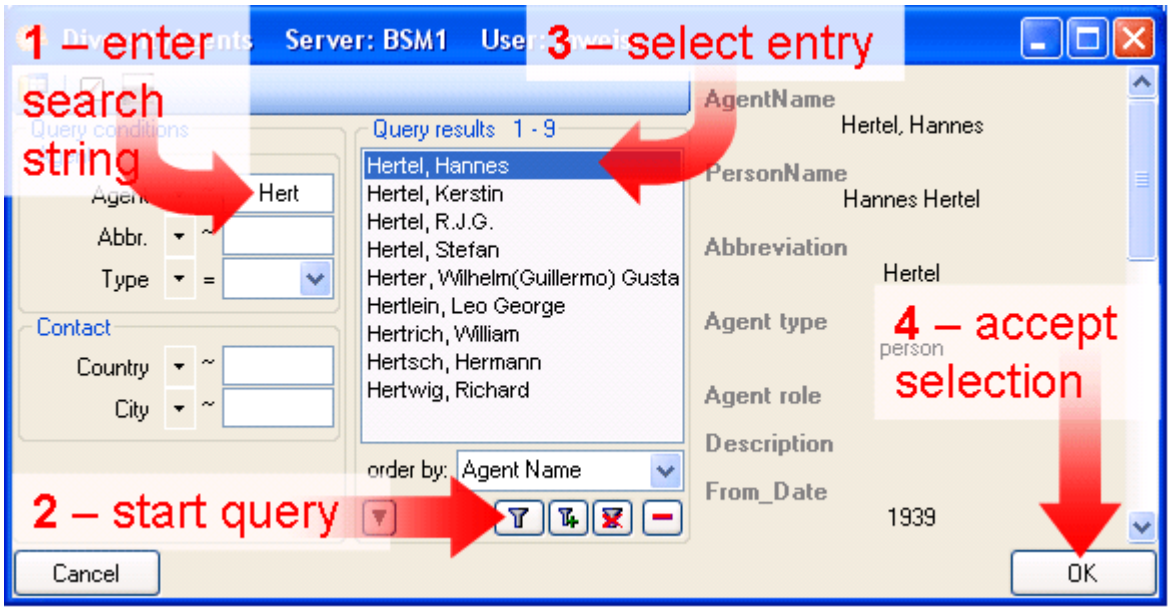


Collectors

Now we will add the collectors of the specimen. In the overview, select the specimen (see point **1** above) then click on the button to insert a new collector (see point **5** in image below). Select the collector in the overview (see point **6** in image below). Now we use the module DiversityAgents to search for a certain person. Click on the button to open the interface to the module (see point **7** in image below).



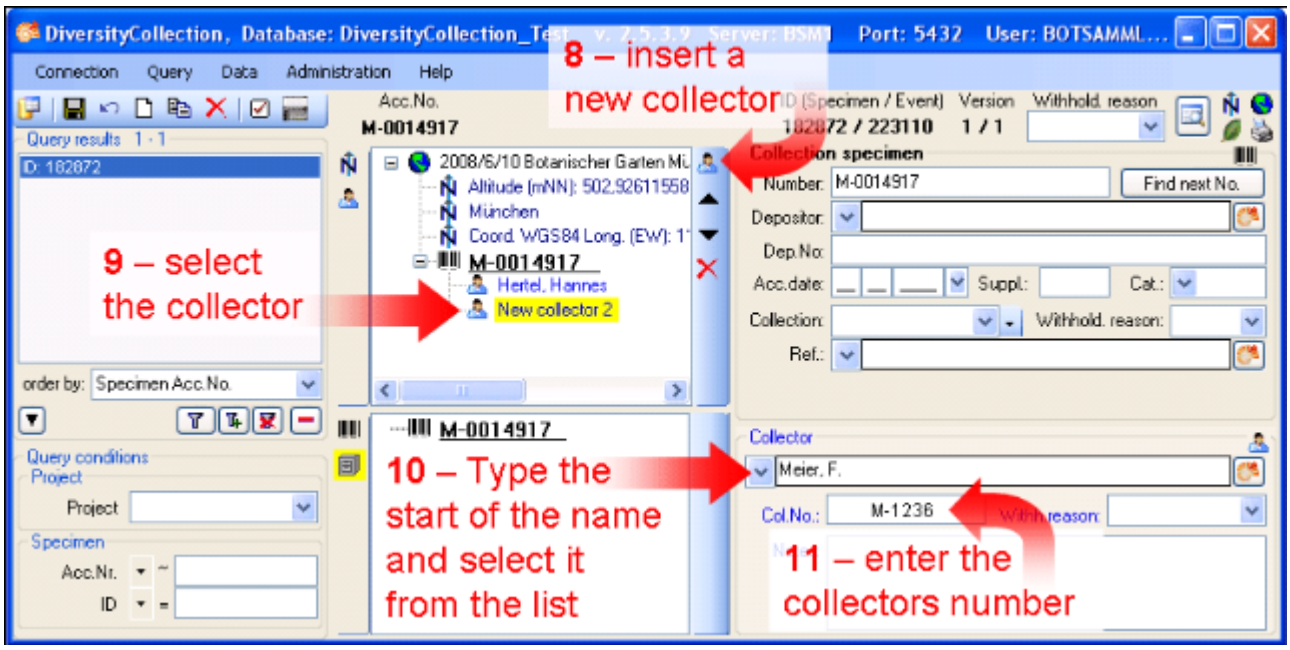
In the interface for DiversityAgents, enter search criteria (see point **1** in image below) and click on the button to start the query (see point **2** in image below).



Select the correct entry from the query result (see point **3** in image above) and click on the OK button (see point **4** in image above) to store the name in DiversityCollection.

The next collector will have no connection to the module DiversityAgents. To insert this collector, again click on the button (see point **8** in image below), select it (see point **9** in image below), type the start of the name in the field (in this example "Mei" would be a good

choice) for the name and click on the drop down button (see point **10** in image below) to select a name from the list of collectors already stored in the database. Finally if the collector has a field number, enter this field number of this collector (see point **11** in image below).



This tutorial is continued in the sections listed below:

- section Collection specimen relations
- section Organisms and identifications
- section Specimen parts and storage

Tutorial - relations between specimen

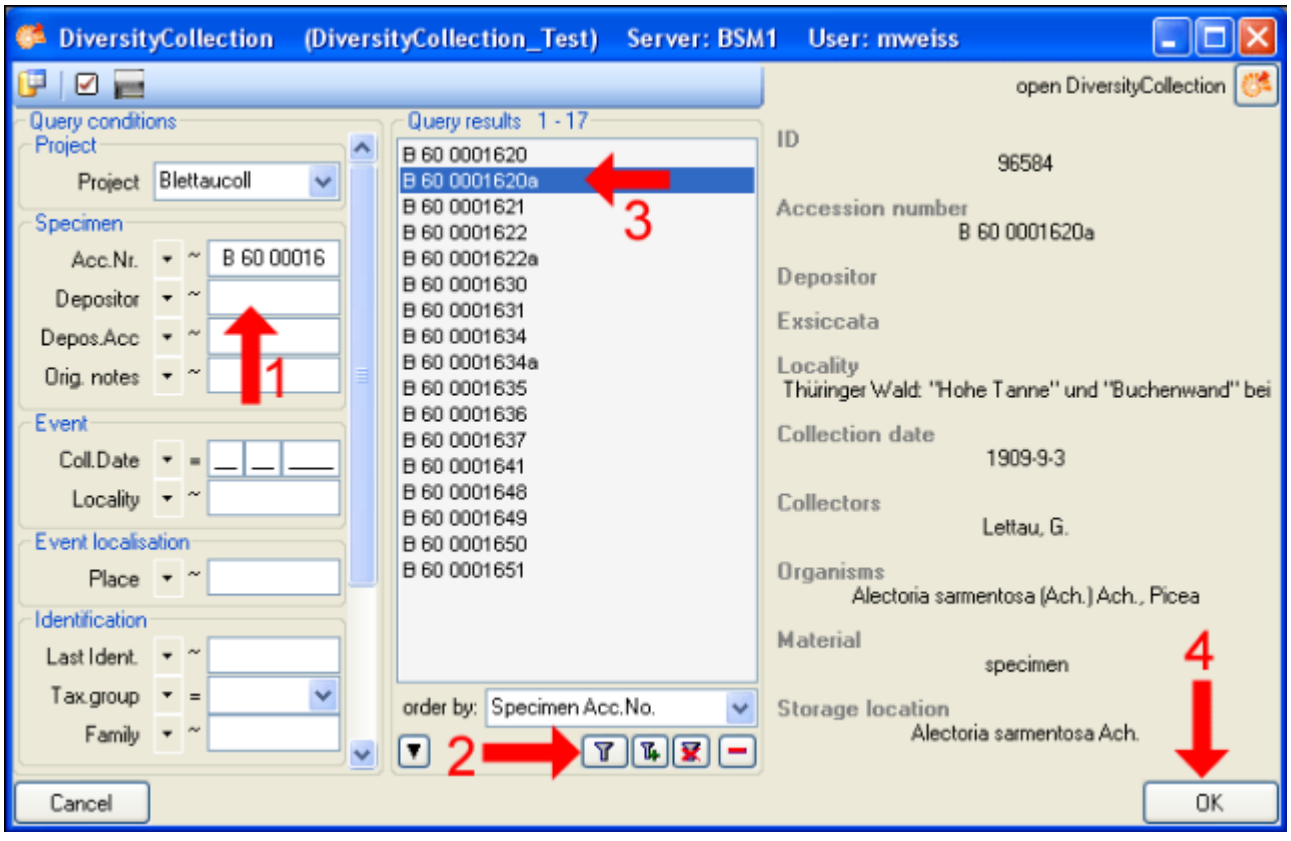
There are two types of relations possible:

1. relation to a specimen within DiversityCollection
2. relation to a specimen not administrated in the local database.

1. Internal relations

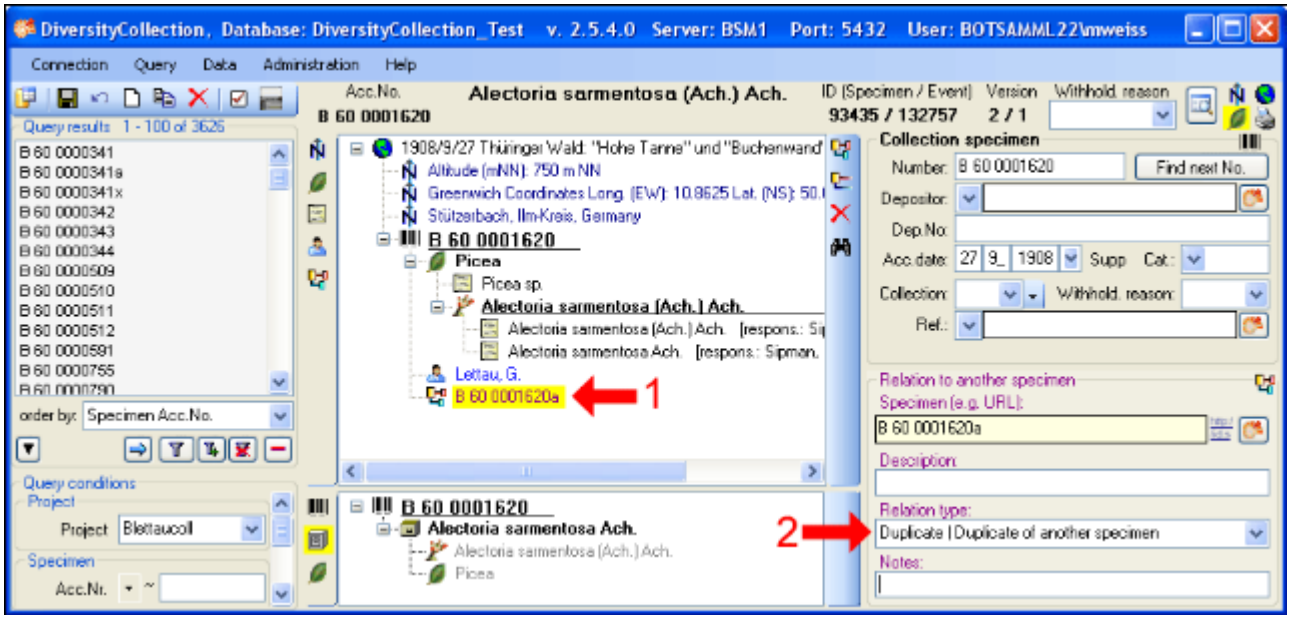
To enter a relation to a specimen in DiverisityCollection select the specimen in the tree (see point **1** in image below) and click on the button (see point 2 in image below).

A window will open where you can search for the related specimen. In the window enter your restrictions (see point **1** in image below), start the query (see point **2** in image below), select the related specimen from the result list (see point **3** in image below) and click OK (see point **4** in image below) to insert the relation.



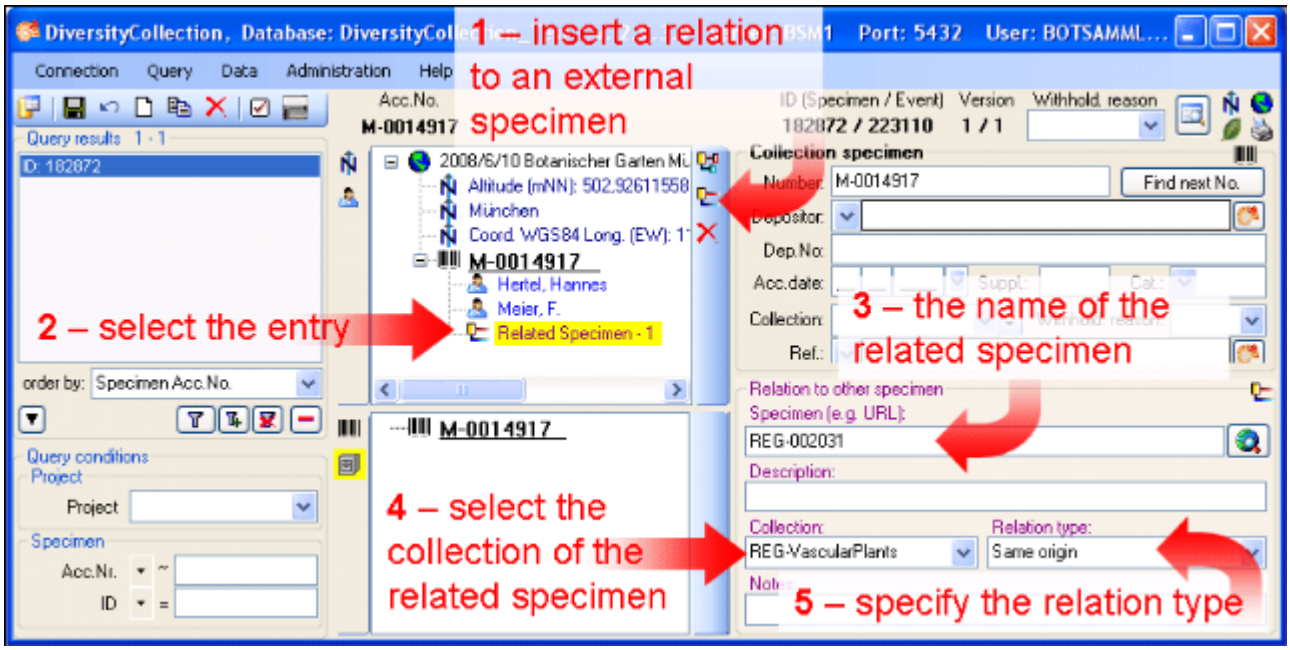
In the window select the entry for the relation (see point **1** in image below) to display the

fields for the details. Then enter the type of the relation (see point **2** in image below).



2. External relations

To enter a relation to a specimen in a foreign collection, insert a relation (see point **1** in image below) and select it in the overview (see point **2** in image below). Enter the name of the specimen (see point **3** in image below) and the type of the relation (see point **5** in image below). .

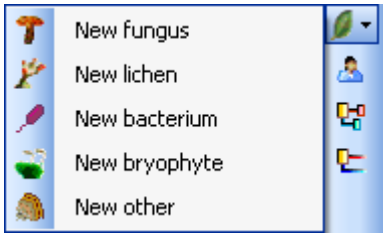


If there is a dataset for the collection available you can select it from the list (see point **4** in image above). Otherwise you first have to add this collection to the list. If you have the permissions to edit the collections choose Administration -> Collections from the menu to add a collection. See the section collection for details.

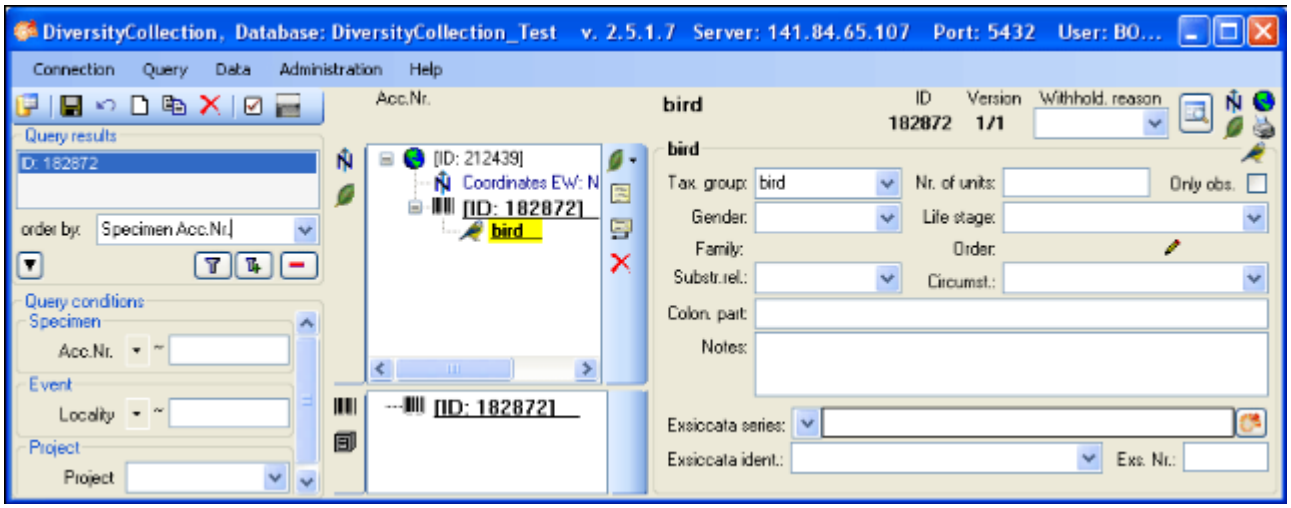
For further informations about the relations turn to the section relation.

Tutorial - organisms and identifications

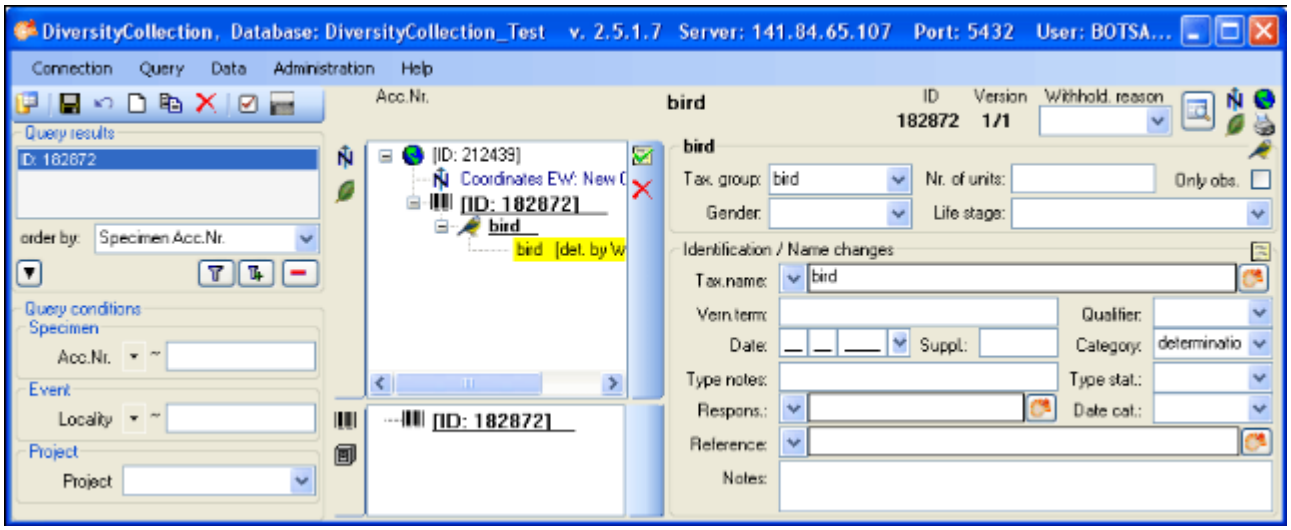
To enter the organism, choose the specimen entry in the tree. Then select the organism from the list.



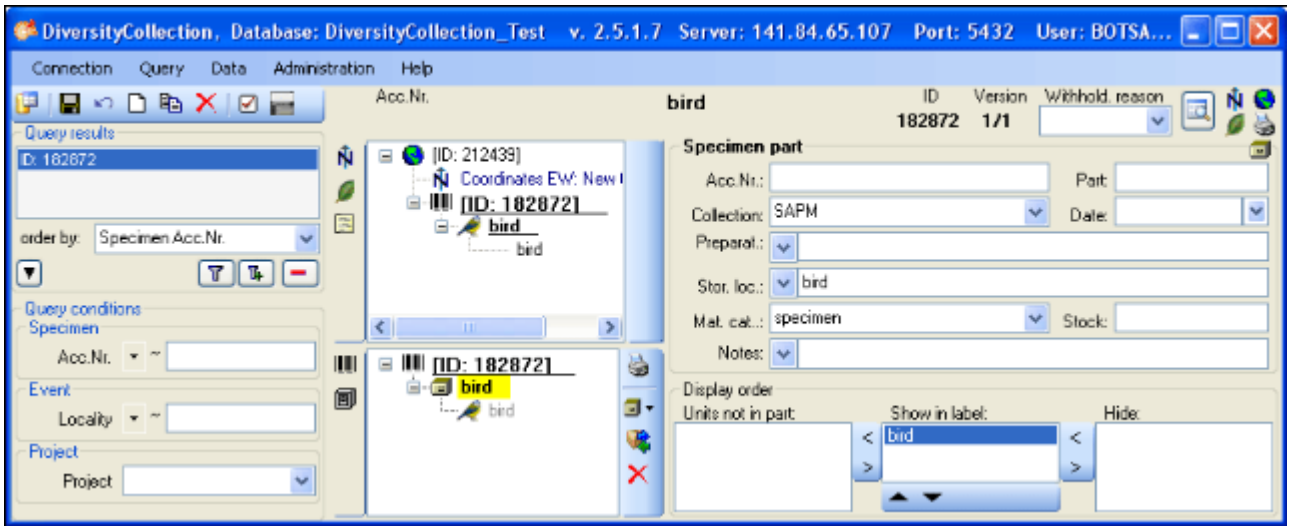
This will enter an entry for the organism unterneath the entry of the specimen as shown below. To enter details for this organism, select the entry in the tree.



To enter an identification, click on the buttom. This will insert a dataset for the idenfication of the organism underneath the organism as shown below. Select this entry to enter the details of the identification.



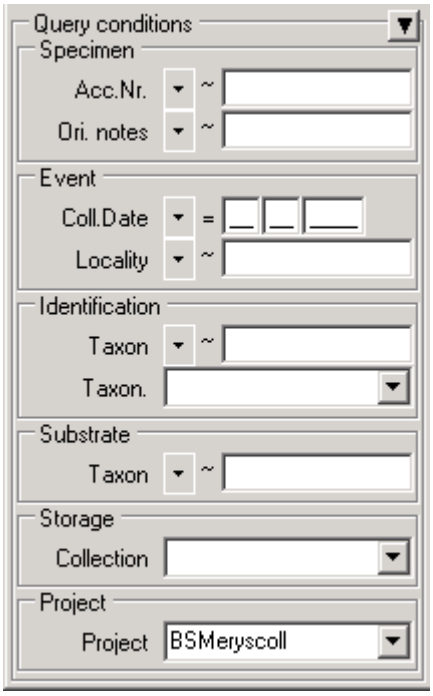
As a last step, enter the data connected with the storage of the specimen. In the bottom tree, select the entry for the specimen and than select a material category from the list. This will enter a dataset for a specimen part underneath the specimen as shown below. Select this entry to edit the data for the collection, the storage location etc.


DiversityCollection, Database: DiversityCollection_Test v. 2.5.1.7 Server: 141.84.65.107 Port: 5432 User: BOTSA...
Connection Query Data Administration Help
Query results
D: 182872
order by: Specimen.Acc.Nr.
Query conditions
Specimen
Acc.Nr.
Event
Locality
Project
Project
Acc.Nr.
bird
ID
182872
Version
1/1
Withhold. reason
[ID: 212439]
Coordinates EW: New
[ID: 182872]
bird
bird
[ID: 182872]
bird
bird
Specimen part
Acc.Nr.:
Part:
Collection:
SAPM
Date:
Preparat.:
Stor. loc.:
bird
Mat. cat.:
specimen
Stock:
Notes:
Display order
Units not in part:
Show in label:
bird
Hide:

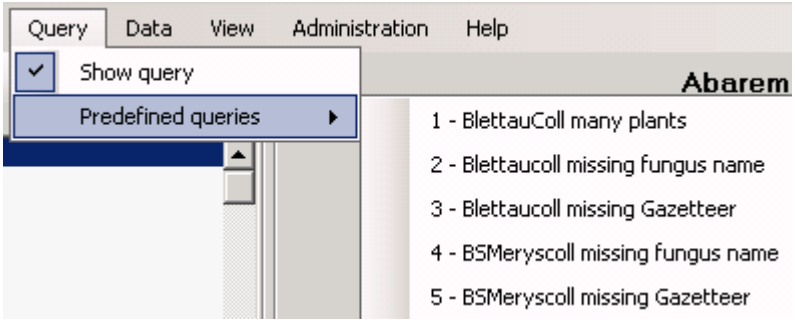
Queries - overview

To search for specimens in the database you can choose 3 options:

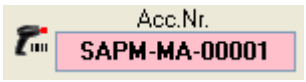
With the user defined queries, you can define any query condition - this is the default query mode.



The predefined queries are defined by the system administrator and are accessible via the menu **Query** - **Predefined queries**. To return to the userdefined click on the **Show query conditions** button.

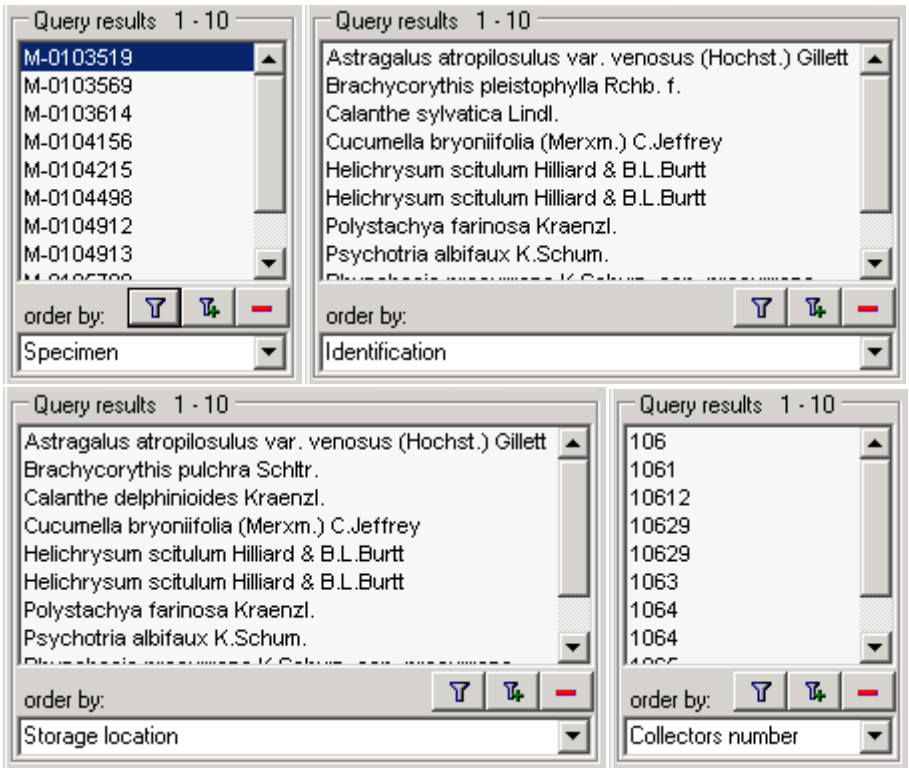


With the scan mode you can use a barcode scanner to search for a specimen. To work with the scan mode, select the **Scan mode** in the **Query** menu. To return to another query mode, deselect the Scan mode.

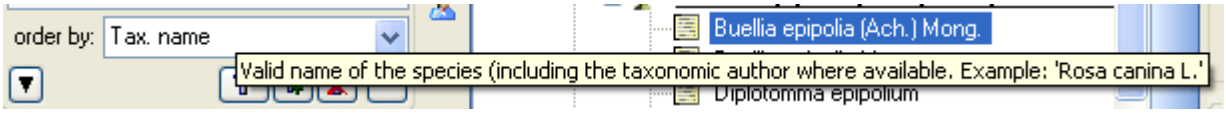


Result list

The result list displays the specimens found in a query.

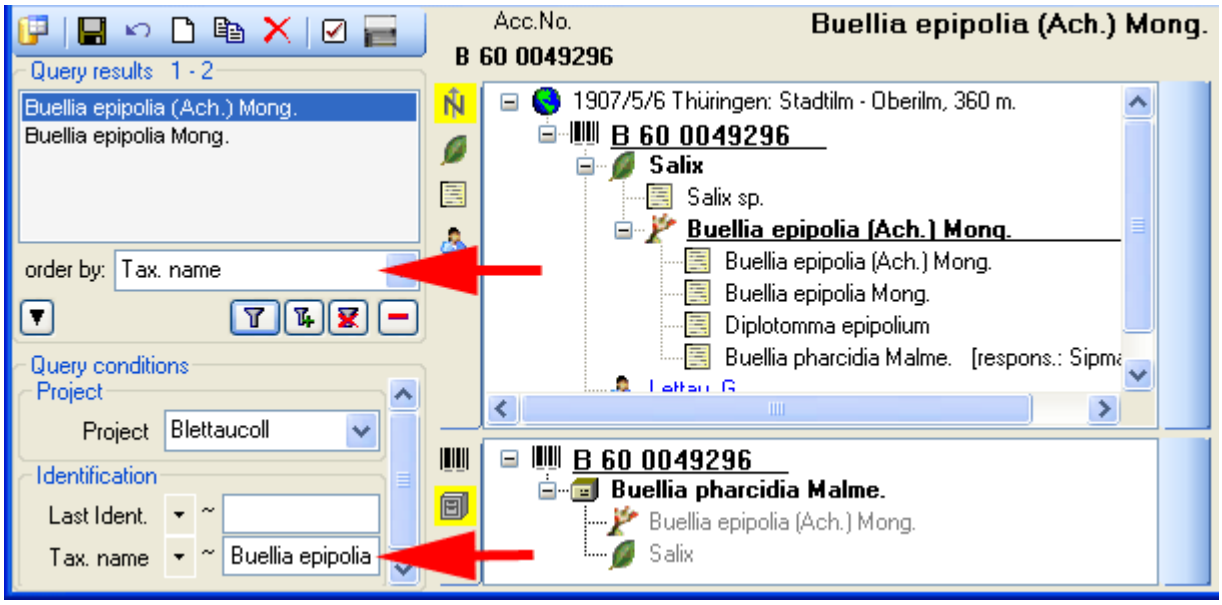


The specimens can for example be shown with their accession number, their identifications or their storage location etc. as shown in the images above. To get further informations about the chosen field, just place the mouse in the field. A text box will appear with the description of the field (see below).

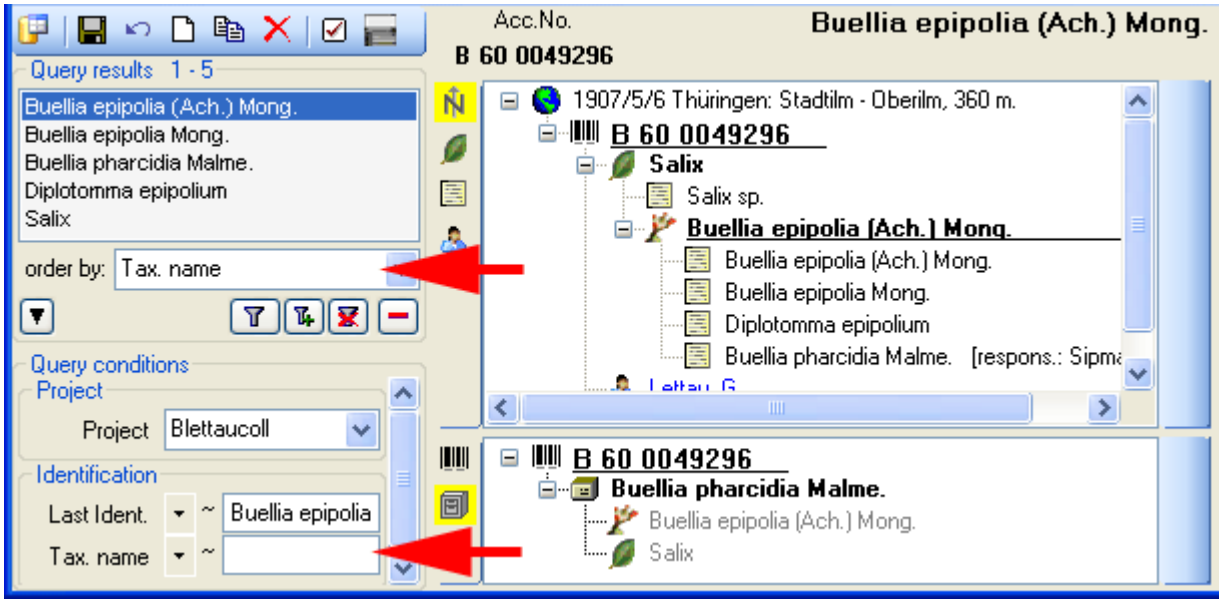


You can restrict the maximal number of specimens together with the query options (click on the button), for example if you have a slow connection to the database. As a default the maximal number is set to 100. If the number of datasets according to your query is higher than the maximal value set in the query options this will be indicated in the header of the list.

To ensure, that restrictions set in the query conditions will be applied to the specimen list make sure that you choose matching restictions and order columns as shown below. In the upper example corresponding fields where used for restricting the query and the display (**Tax. name**). Here the Query results will be restricted to this field.



In the second example a different field for the restiction was chosen (**Last ident.** <> **Tax. name**). The query result in consequence will list all entries found in the field **Tax. name** from the datasets where on entry matches the restriction (see below).



To search for specimens, enter the restrictions in the fields for the search conditions and click on the button. The specimens found in the database will be shown in the result list. To add specimens with differing search conditions click on the button. If the list of items is longer than your maximal number of returned items you can browse the next items with the button. If you want to remove entries from the selected list, choose them in the list and click on the button. This will not delete the data from the database, but remove them from your query result.

Here some examples you can select for display in the result list:

AccessionNumber: One entry is shown for each specimen with its corresponding accession number.

Last identification: The last identification for every unit in a specimen is shown in the list. As

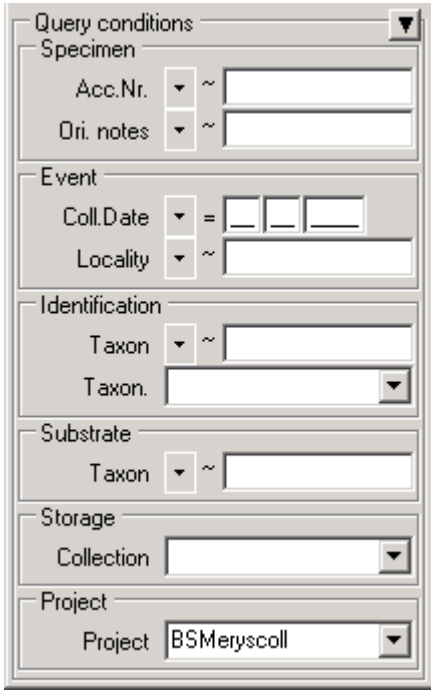
there can be several units in one specimen several entries for one specimen may appear in the list.

Storage location: The storage location of every part of a specimen stored in the collections is shown in the list. As parts of a specimen can be stored in several collections under different names several entries for one specimen may appear in the list.

Collecting number: The collecting number given by the collector of every sample of a specimen is shown in the list. A specimen can have several collectors each with different number. Therefore several entries for one collection specimen may appear in the list.

Query

There are two ways to search for specimens in a collection. The options for a fast search are displayed in the main window beneath the list of the items. You can change this arrangement using the / button to place the query options on the left side of the item list.



To search for specimens enter the restrictions in the fields for the search conditions and click on the button. The specimens found in the database will be shown in the specimen list. To add specimens with differing search conditions click on the button. To clear all entries in the query fields use the button. If the list of items is longer than your maximal number of returned items you can browse the next items with the button. To move back to the previous block of items click on the button. If you want to remove entries from the selected list, choose them and click on the button. This will not delete the data from the database, but remove them from your query result.

Within the query options you have several possibilities to specify your search restriction. Use the drop down button to change between the operator. The available operators are shown in the table below.

Operator	Meaning	Example
Text		
~	search for an entry like ...	Pinus s[iy]lvestris % (you can use wildcards)
=	search for an entry exactly equal to ...	Pinus silvestris L.
≠	search for an entry not like ...	Pinus s[iy]lvestris % (you can use wildcards)

∅	search for an entry where a value is missing ...	
•	search for an entry where a value is present ...	
-	search for an entry between ... and ...	2000 - 2005
Numeric		
=	search for an entry exactly equal to ...	2006
<	search for an entry lower than ...	2006
>	search for an entry bigger than ...	2006
-	search for an entry between ... and ...	2000 - 2005
∅	search for an entry where a value is missing ...	
•	search for an entry where a value is present ...	
Date		
=	search for an entry exactly equal to ...	20.3.2006
<	search for an entry lower than ...	20.3.2006
>	search for an entry bigger than ...	20.3.2006
Hierachy		
=	search for an entry exactly equal to ...	M-Fungi
≠	search for an entry that is not equal to ...	M-Fungi
∅	search for missing entry ...	M-Fungi
•	search for present entry ...	M-Fungi
Δ	search including childs in a hierarchy ...	M-Fungi

To hide the area containing the search fields click on the button. If the search area is hidden and you want to start a new search, just click on the button.

To change the displayed fields for searching specimens click on the button to change the query options.

Scan mode

To search for a specimen with the help of a barcode-scanner select the **Scan mode** from the Query menu. The query part will be hidden and the field for the accession number will then be accessible for the entry with the scanner. If the field for the entry of the accession number Acc.Nr.: is not activated, move the mouse to the field to activate it. Then scan the barcode and the program will start the search for the specimen in the database.

Grid mode

To edit the data in a data grid, choose the **Grid mode** from the Query menu. The query part will be hidden and table will appear where every dataset from the query result list is

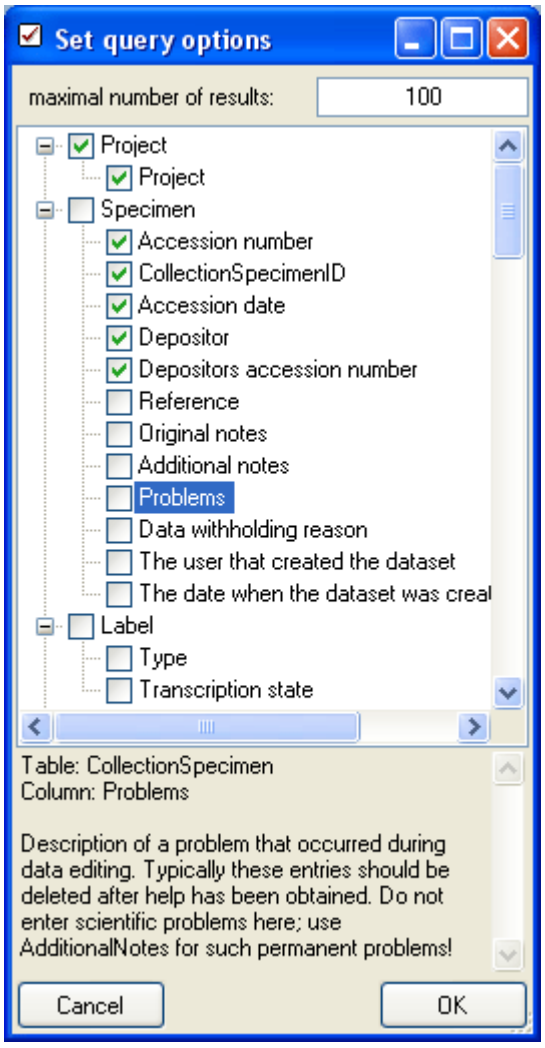
restricted to one line. Please keep in mind, that in this view, you can only see a limited part of the data. So for example you will only the the last identifications for an organism. The selection of the visible fields can be adapted in the tree above the list.

To replace a part of a text in a column, select the column, enter the text that should be replaced and the replacement in the corresponding fields (see below). To start the replacement click the button.

Query options

The maximal number of items shown in a query result can be set in the window for the query options. The default value is set to 100. If you have a fast connection to your database or need to see more or less results, you may change this value to any number you like.

To change the displayed search fields click on the button. This opens a form where you can select and deselect the fields shown for searching specimens. You might also change the maximum number of items that will be shown in the result list.



After having edited the query options click OK to store you selection. The new selection will become active for the next query.

Wildcards in SQL

There are 4 different possibilities for wildcards in SQL:

% any string consisting of no, one or many characters, e.g. Pinus **%** will find anything like Pinus, Pinus sylvestris, Pinus strobus etc.

***** any string consisting of no, one or many characters, e.g. Pinus ***** will find anything like Pinus, Pinus sylvestris, Pinus strobus etc.

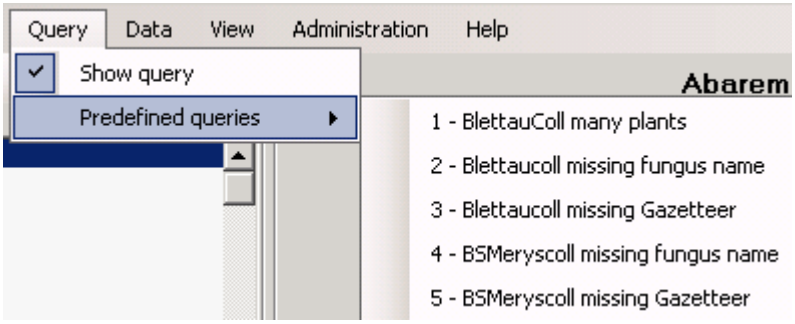
_ a single character, e.g. Pinus s**_**lvestris will find Pinus s**y**lvestris and Pinus s**i**lvestris etc.

[] any character out of a given range like **[abcde]** or **[a-e]**, e.g. Pinus s**[iy]**lvestris will find Pinus sylvestris and Pinus silvestris.

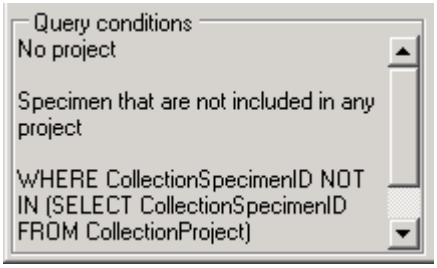
[^] any character not in a given range like **[^abcde]** or **[^a-e]**, e.g. Pinus s**[^i]**lvestris will find Pinus sylvestris but not Pinus silvestris.

Predefined queries

Besides setting querries for specimens via the query options you can define separate predefined user-specific queries. These are listed under the menu topic **Query - Predefined queries**.

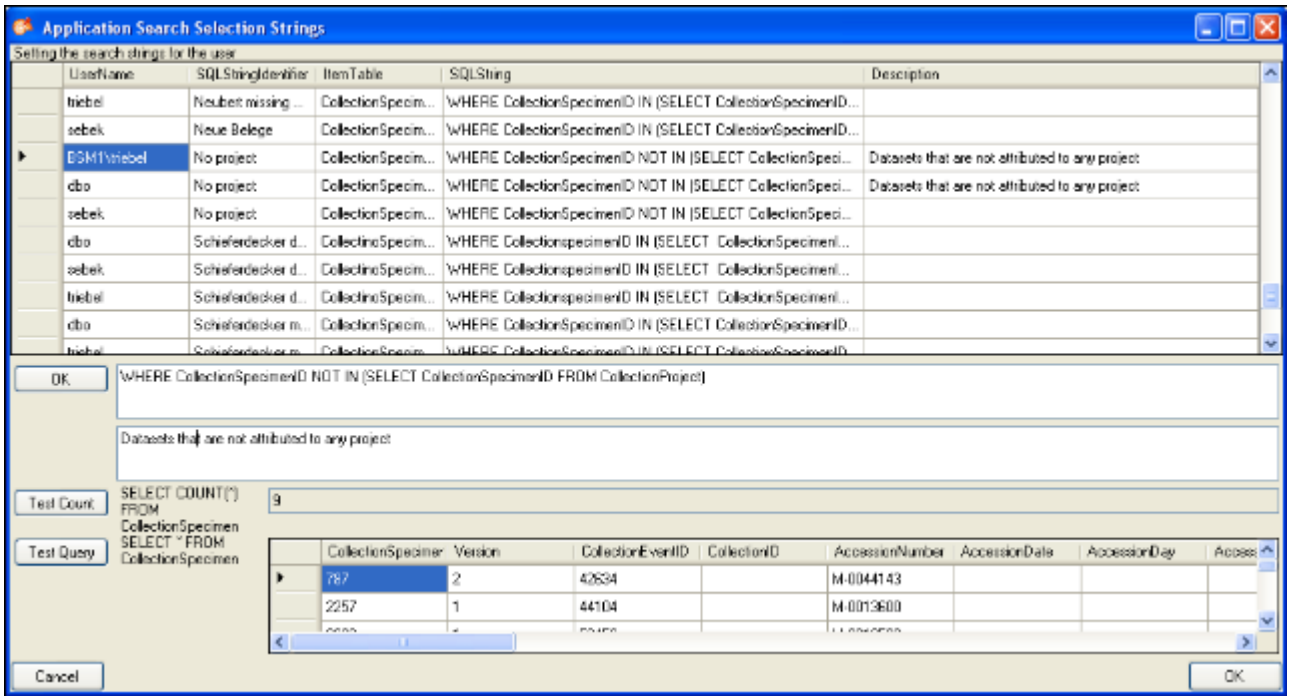


If you choose one of these predefined queries, the query options will be hidden and the command of the query will be shown at the base of the specimen list. The first line shows the title of the query, the next lines contain the description followed by the part of the query command that restricts the selection of the datasets (= WHERE-clause of the SQL-statement).



To return to the user defined query click on the **Show query conditions** button.

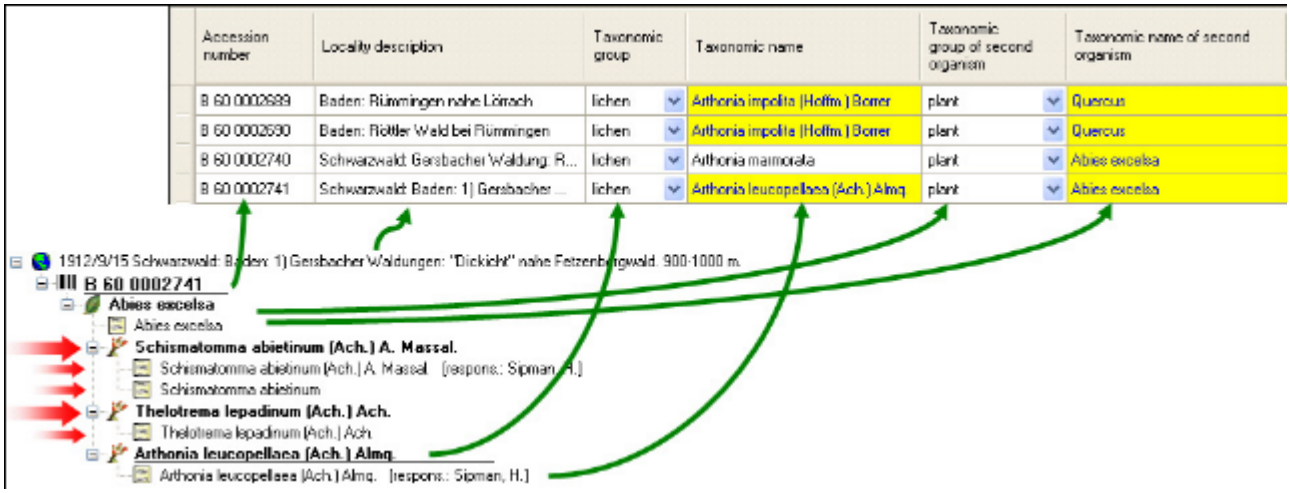
If you are an administrator you can create new queries for users. To create a predefined query choose **Administration - Queries...** from the menu. A window as shown below will open, where you can create, edit and test your queries.



In the upper field you define the WHERE-clause of the SQL string of your query. Keep in mind that the queries can refer to different tables, depending on the order column chosen by the user. So queries in DiversityCollection should start with the reference to the primary key of the main table (CollectionSpecimenID in table CollectionSpecimen and depending tables). The lower field contains the description for the query as shown in the user interface. To test a query use the **[Test count]** and **[Test Query]** buttons.

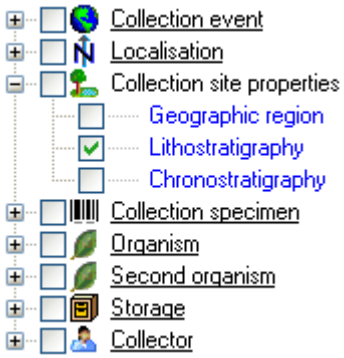
Grid mode

To edit the data in a data grid, choose the **Grid mode** from the **Query** menu. A window will open where every dataset from the query result list is restricted to one line. Please keep in mind, that in this view, you can only see a limited part of the data. So for example you will only see the last identification of an organism. This is demonstrated in the image below, where two organsims, indicated with the red arrows will not appear in the grid.



Customize visibility of fields

The selection of the visible fields can be adapted in the tree above the list. Change the selection of the colums and click on the **[Set columns]** button.

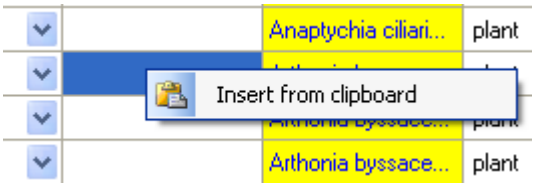


Find and replace

To edit the values in a column with e.g. the replace function, select a field in that column. If only a part of the fields should be edited, select these fields. Otherwise all fields in the column will be edited. Then choose the function you want to apply (remove, insert, append or replace). To replace a part of a text in a column, enter the text that should be replaced and the replacement in the corresponding fields. To start the **replacement** click the button. To insert a string to the **beginning** of all entries in a column, click the button. To **append** a string to all entries in a column, click the button. To **remove** all entries from a column click the button.

Transfer from spreadsheet

You can transfer data from a spreadsheet, e.g. Excel or Calc. Copy one column of these data in the spreadsheet and then in DiversityCollection, click in the upmost cell where these data should be inserted and use the context menu (click the right mouse button) to insert the data.



Editing

Some columns can not be edited directly but are linked to external modules or services. These columns appear as buttons. Just click on the button the call the service. If a value is linked to an entry in an external module, the background will change to yellow and you can not change the text. Some values are linked to a list of values. Use the drop down list to change the value in one of these columns.

If you click in the empty line at the base of the data grid, you will be asked if you want to create a new dataset. The programm will ask you for a new accession number and the project of the new dataset. Another way to create a new dataset is the copy button . Just click in the line you want to create a copy of and then click on the copy button . A copy of the dataset will be inserted at the base of the datagird.

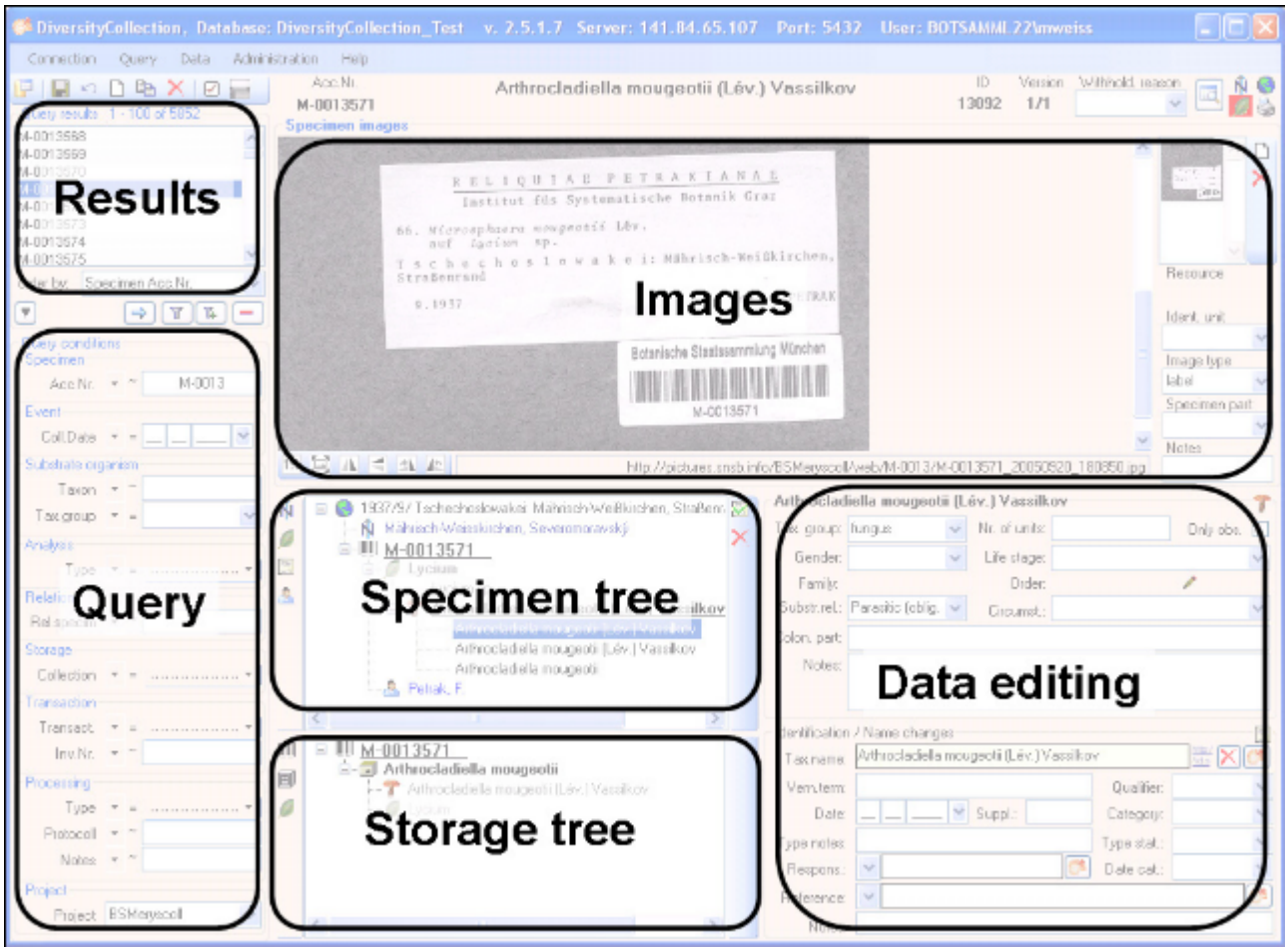
Saving the data

To save all changes click on the button. To undo the all changes since the last time the datasets were saved, click on the button. To save the changes in the current dataset, use the button. To undo the changes in the current dataset, click the button. If you click the **[OK]** button, you will be asked if you want to save the changes before the window will be closed. If you click the **[Cancel]** button or close the window your changes will not be saved. To save the data shown in the grid in a tab delimited file, click on the button.

To change to a selected dataset in the main form, click the button.

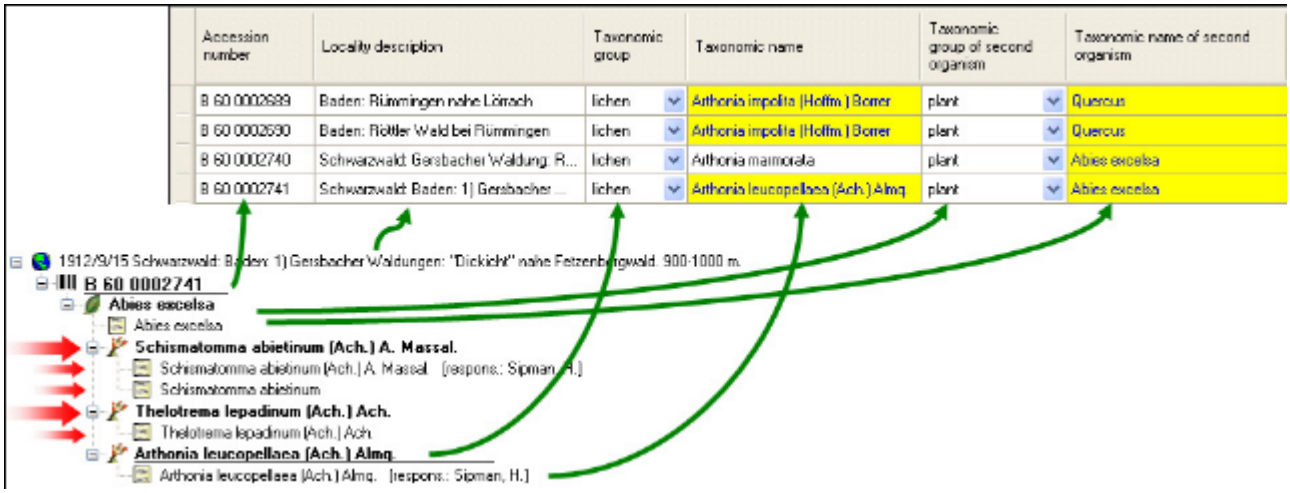
Editing the data

The main window of the DiversityCollection client contains two main areas. At the left you find the query and the results of this query. In the right part the data of the dataset selected in the result list is shown. The upper part of the data area shows the images, labels etc. In the lower part you find two trees that give you an overview and access to the data. The data of an entry selected in one of the trees are shown in the data editing part.



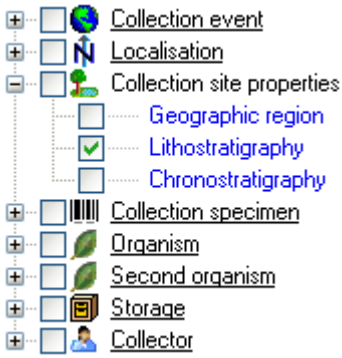
Grid mode

To edit the data in a data grid, choose the **Grid mode** from the **Query** menu. A window will open where every dataset from the query result list is restricted to one line. Please keep in mind, that in this view, you can only see a limited part of the data. So for example you will only see the last identification of an organism. This is demonstrated in the image below, where two organsims, indicated with the red arrows will not appear in the grid.



Customize visibility of fields

The selection of the visible fields can be adapted in the tree above the list. Change the selection of the colums and click on the **[Set columns]** button.

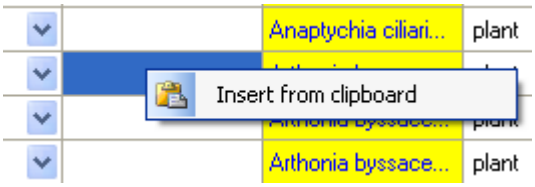


Find and replace

To edit the values in a column with e.g. the replace function, select a field in that column. If only a part of the fields should be edited, select these fields. Otherwise all fields in the column will be edited. Then choose the function you want to apply (remove, insert, append or replace). To replace a part of a text in a column, enter the text that should be replaced and the replacement in the corresponding fields. To start the **replacement** click the button. To insert a string to the **beginning** of all entries in a column, click the button. To **append** a string to all entries in a column, click the button. To **remove** all entries from a column click the button.

Transfer from spreadsheet

You can transfer data from a spreadsheet, e.g. Excel or Calc. Copy one column of these data in the spreadsheet and then in DiversityCollection, click in the upmost cell where these data should be inserted and use the context menu (click the right mouse button) to insert the data.



Editing

Some columns can not be edited directly but are linked to external modules or services. These columns appear as buttons. Just click on the button the call the service. If a value is linked to an entry in an external module, the background will change to yellow and you can not change the text. Some values are linked to a list of values. Use the drop down list to change the value in one of these columns.

If you click in the empty line at the base of the data grid, you will be asked if you want to create a new dataset. The programm will ask you for a new accession number and the project of the new dataset. Another way to create a new dataset is the copy button . Just click in the line you want to create a copy of and then click on the copy button . A copy of the dataset will be inserted at the base of the datagird.

Saving the data

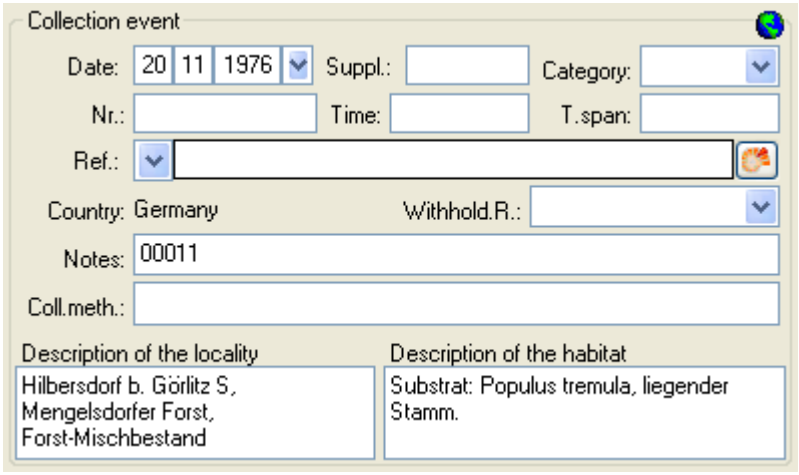
To save all changes click on the button. To undo the all changes since the last time the datasets were saved, click on the button. To save the changes in the current dataset, use the button. To undo the changes in the current dataset, click the button. If you click the **[OK]** button, you will be asked if you want to save the changes before the window will be closed. If you click the **[Cancel]** button or close the window your changes will not be saved. To save the data shown in the grid in a tab delimited file, click on the button.

To change to a selected dataset in the main form, click the button.

Collection event

Specimens stored in a collection are gathered during a collection event. This collection event keeps information about the geographic locality, the habitat, the collection date etc. During an collection event, several specimens may have been collected. If a specimen is not assigned to a collection event, you can assign the specimens to an existing event with a click on the button. To create a new event for a specimen click on the button. If you assign the specimen to an existing event, a window will open as shown below where you can search for the events already included in the database. Choose an event and click OK to assign the specimen to this event. In the tree view, the collection event is symbolized with an icon as shown below.

To edit the data of the collection event, choose it in the tree view to open the detail fields as shown below. If the collection date does not correspond to a certain day, you may use the **Suppl.** field to enter e.g. a range or a series of dates or the **T.span** field to document a certain timespan.



The text shown in the tree view is composed of the date of the collection event and the description of the locality. For each collection event you can enter several geographical localisations and properties. To see the locality according to the coordinates stored in the database you can check the Maps. For each collection event you can enter images related to this event.

If other specimens were collected during the same collection event, this will be visible if you show the whole hierarchy of the event series. To do this click on the button in the panel on the left of the tree. See event series for further details. Here you can move a specimen to another event by drag and drop.

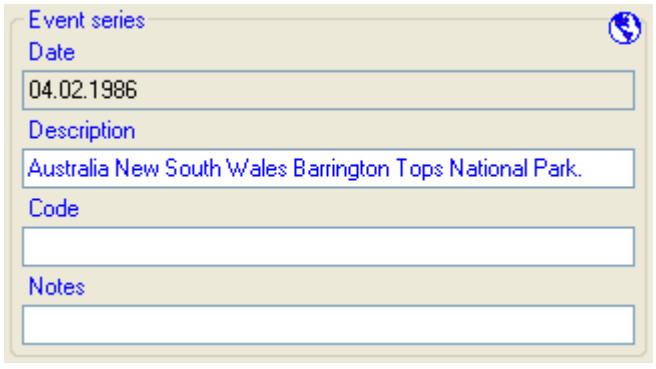
Data are stored in the table CollectionEvent.

Collection event series

If you need a hierarchical order of your collection events respectively to organise your collection events e.g. to document expeditions, you can do this with a collection event series. For a better differentation between events and collection event series you have a blue text in the hierarchy and the editing part and a different icon . A collection event series can contain other collection event series and collection events. Information about the geographic locality, properties of the collection site, the date of collecting etc. are stored in the collection event. To show or hide the collection event series you have two options. In the panel of the right of the tree the button will show the superior event series of the current collection event as shown below.

The button will show the whole hierarchy of the event series as shown below.

To edit the data of an event series, select it in the tree to display the fields with the details as shown below.



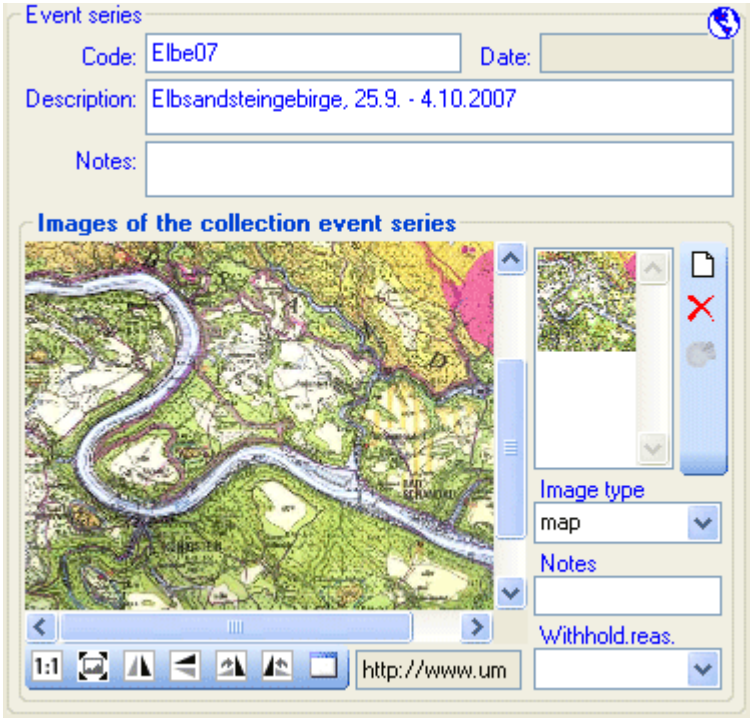
To insert a new collection event series click on the button. If there are no collection event series so far, the collection event will be placed within the new collection event series. If there are collection event series present, the new collection event series will be placed below the selected collection event series. To assign an collection event to an existing collection event series already available in the database click on the icon.

To move an item within the hierarchy, just drag it with the mouse to whatever position it should be placed. Keep in mind, that specimens can only be placed in collection events and collection events only in collection event series.

If you want to delete a collection event series or an collection event, remove all depending collection event series, events and specimens and click on the button. A specimen can not be deleted here. If you want to remove a collection event from a collection event series, click

on the button to open the window for selecting a collection event series. Then select nothing but simply click OK to remove the link to the collection event series.

The images for a collection event series are shown below the data of the collection event series as shown below.To add images to a collection event series, click on the button, to remove an image use the button.



To zoom a sector of the image, just drag the mouse over the image. A red square will indicate the zoomed area. To set the size of the image to the original resolution click on the button. To adapt the size of the image to the available space in the form click on the button. To change the orientation of the image use the appropriate buttons (flip horizontal, flip vertical, rotate right, rotate left). To view the image in a separate form, click on the button. If the image should not be published e.g. on the internet, enter any reason in the **Withhold. reason** - field.

If you want to change to another specimen listed in the collection event series select it in the hierarchy and click on the button.

The current specimen together with the event and all superior collection event series will be highlighted.

The data about the collection event series are stored in the table CollectionEventSeries.

Event images

The specimens stored in a collection are gathered during a collection event. To insert an image related to a collection event click on the button. If you want to delete an image, click on the button.

If for any reason, an image should not be published i.e. shown on a website, enter the reason in the field **Withholding reason**. Only images where this field is empty will be shown e.g. on websites.

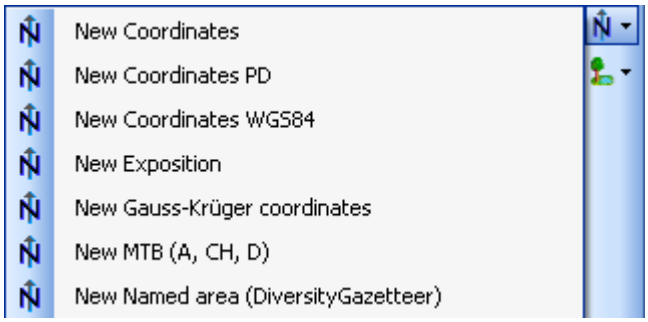
To zoom a sector of the image, just drag the mouse over the image. A red square will indicate the zoomed area. To set the size of the image to the original resolution click on the button. To adapt the size of the image to the available space in the form click on the button. To change the orientation of the image use the appropriate buttons (flip horizontal, flip vertical, rotate right, rotate left). To view the image in a separate form, click on the button. If the image should not be published e.g. on the internet, enter any reason in the **Withhold. reason** - field.

Data are stored in the table CollectionEventImage.

Geography

As additional information to the description of the locality you can use several localisation systems, e.g. georefencing. These entries are marked with an icon in the tree as shown below.

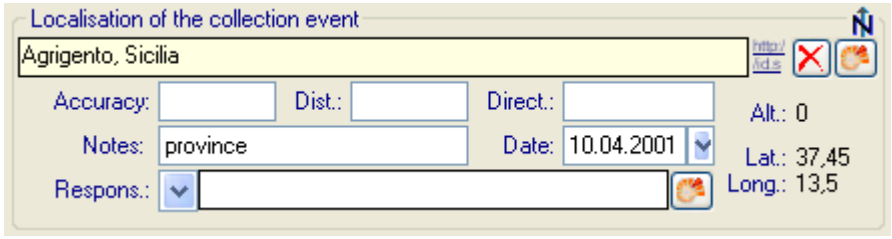
To add a new entry choose the localisation system from the drop down list as shown below. Only items that are not already set for a collection event will be shown. The items that are visible in the dropdown list can be customized.



If a localisation should be removed, select it from the list and click on the button.

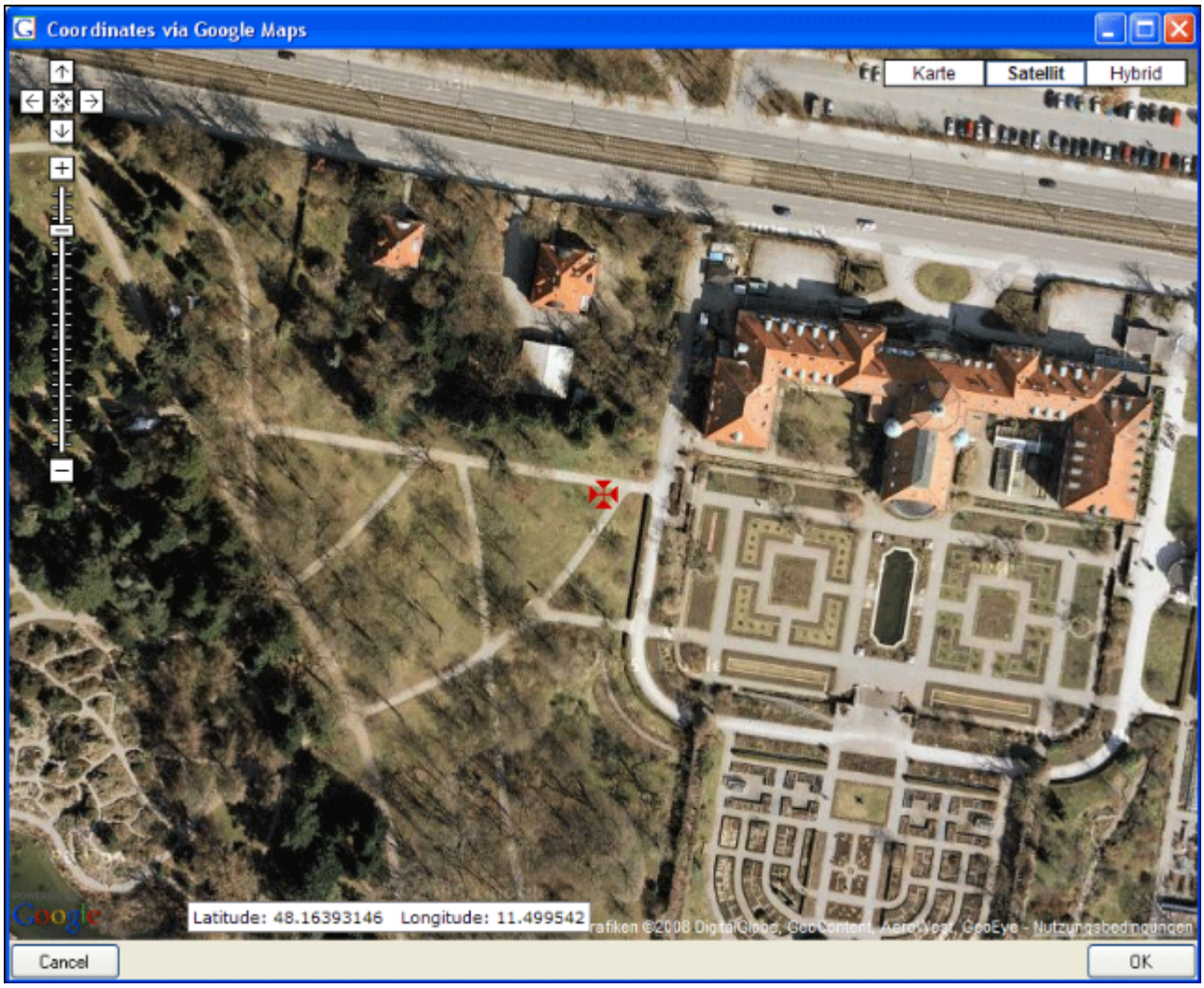
Named areas - DiversityGazetter

To edit the details of an entry, select in the list to open the fields in the form on the right. The localisation system DiversityGazetteer is linked to the module DiversityGazetteer within the Diversity Workbench, providing information on geographical names as shown below.



Coordinates (WGS84)

If you use geo-coordinates as georeferencing system (coordinates WGS84), a button will appear, that will provide you with the possibility to set or correct the coordinates via Google Maps. Just click on the button, and a window will open where you will see a map provided by Google Maps as shown below. If there are allready coordinates provided by a different localisation e.g. by the DiversityGazetteer, these will be taken as a starting position. So you may use an entry for a named area (DiversityGazetteer) as a approximate localisation and than additional coordinates as the more accurate localisation. C

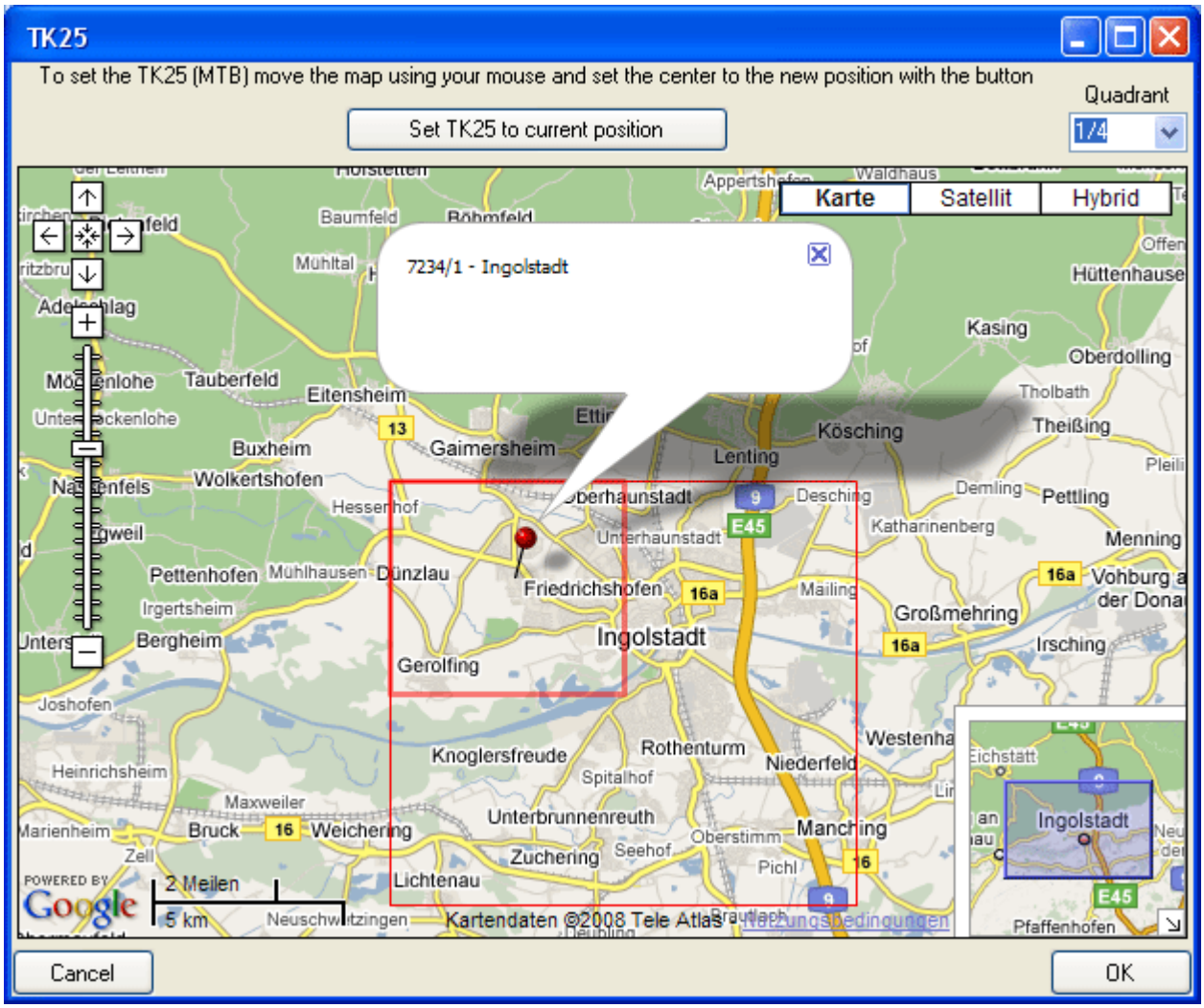


If you start with existing geographic coordinates (WGS 84) in your data, the system will use these as starting point. You can zoom the map, drag it to another position and change from map to satellite mode. The actual position is symbolized by the ✠ symbol in the middle of the map. The current coordinates are shown in the field at the buttom of the map. To take these coordinates in your data, just click OK.

Please keep in mind, that Google Maps coordinates are based on **WGS84**.

TK25 (MTB)

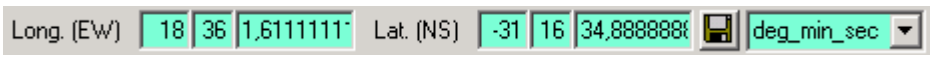
If you choose MTB resp. TK25, the button ✠ will open a window where you can set the TK25 together with the quadrant. See image below. Choose the quadrant, depending on your preferred resolution. The thick line shows the current quadrant, the thin line the whole TK25. The needle in the center is placed at you current position. If you click on it, a messagebox will show you the TK25 informations as shown below.



If you click **[OK]**, the values for the values for the TK25 together with the coordinates of the center of the selected quadrant will be taken into the database.

Conversion of values

The values for any localisation system are stored in two text fields. You can enter your values as simple text. This may prevent any calculations with your values. So you should prefer to enter your values according to the measurement units available. Whatever antique measurement units like feet or Fahrenheit may exist, within the Diversity Workbench measurement data are stored in units according to the *Système international d'unités* (SI). For those who still need to use these units DiversityCollection provides a possibility to convert them into their modern counterparts. To enter a numeric value choose the unit you prefer from the list. The form will change as shown below.



You can then edit the values and click on the button to take the changed values in your data. The system will calculate the corresponding value for storage in the database together with the default accuracy. Correct this accuracy if you have more exact values. To indicate that the shown values are calculated from the values in the database the fields have a green background. When you save the data, DiversityCollection will store geographic coordinates and the average altitude where available in separate fields.

The altitude, the exposition or the slope may be entered as one value or as a range of two

values. If you use the conversion function and have only one value, make sure, that the second field is empty.

Available units: meter and feet.

Available units: Orientiation (N, NE, ...) and degree rel. to North.

Available units: degree and percent.

The accuracy resp. uncertainity will be calculated as an approximation in parts derived from Wieczorek, J. 2001 (MaNIS/HerpNet/ORNIS Georeferencing Guidelines. University of California, Berkeley: Museum of Vertebrate Zoology) and in parts according to Wieczorek, J., Q. Guo, and R. Hijmans 2004 (The point-radius method for georeferencing locality descriptions and calculating associated uncertainty. International Journal of Geographical Information Science 18: 745-767). The unit of the accuracy will always be the unit stored in the database according to the *Système international d'unités* (SI) resp. degrees for angles.

The data for the geography are stored in the table CollectionEventLocalisation.

Maps of the collection event

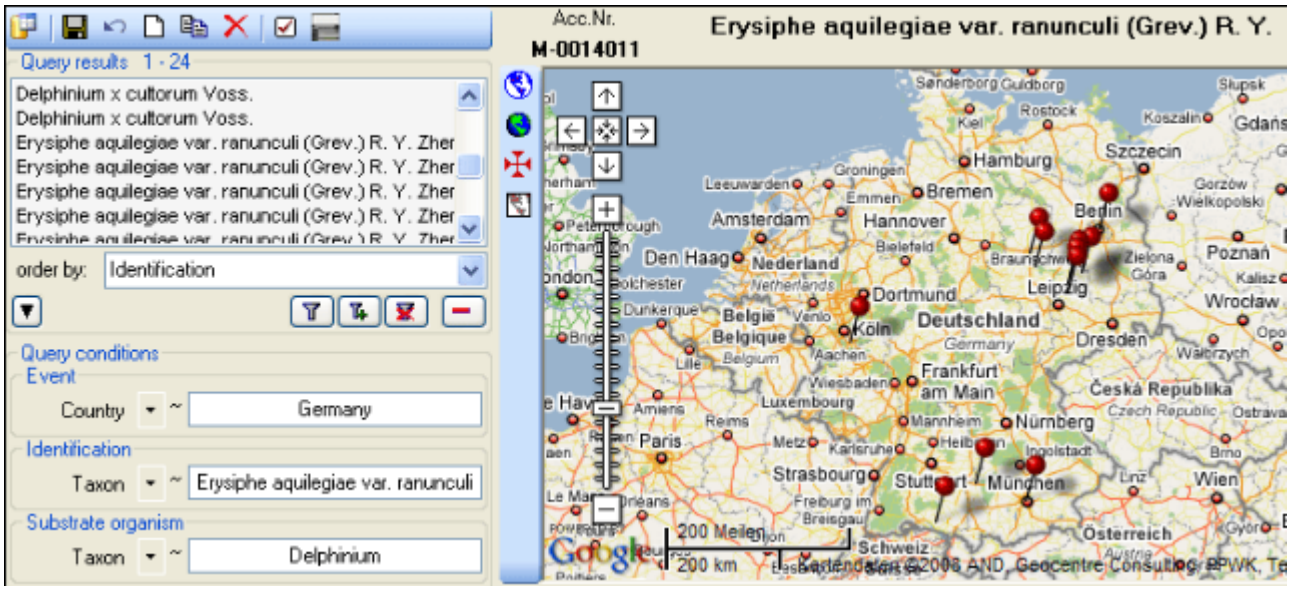
If there are coordinates available for your collection site of your locality using

To use this service, you need access to the internet. Click on the button in the control to show the maps. Click on the button to see the location of your event or if the event is part of a series the button to see all locations of the event series. The site can be displayed either as a satellite image

or a geographical map. If you click on one of the needles marking the locations of the events, a window will show you the description of the event.

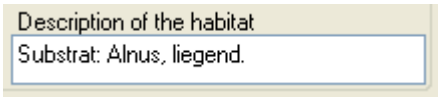
To generate a map you need coordinates stored in the table CollectionEventLocalisation.

To show a distribution map as shown below of all the specimens found in your query, use the button. This functionality is limited in the number of specimens that can be depicted, so it may be neccessary to restrict your query.

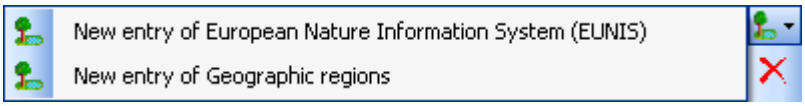

Query results 1 - 24
Delphinium x cultorum Voss.
Delphinium x cultorum Voss.
Erysiphe aquilegiae var. ranunculi (Grev.) R. Y. Zher
Erysiphe aquilegiae var. ranunculi (Grev.) R. Y. Zher
Erysiphe aquilegiae var. ranunculi (Grev.) R. Y. Zher
Erysiphe aquilegiae var. ranunculi (Grev.) R. Y. Zher
order by: Identification
Query conditions
Event
Country ~ Germany
Identification
Taxon ~ Erysiphe aquilegiae var. ranunculi
Substrate organism
Taxon ~ Delphinium
Acc.Nr.
M-0014011
Erysiphe aquilegiae var. ranunculi (Grev.) R. Y.
Hamburg
Bremen
Amsterdam
Hannover
Berlin
Den Haag
Nederland
Dortmund
Deutschland
Leipzig
Dresden
Köln
Frankfurt am Main
Nürnberg
München
Strasbourg
Paris
Wien
Česká Republika
Österreich
Schweiz
200 Meilen
200 km
Google

Habitats and properties of the collection site

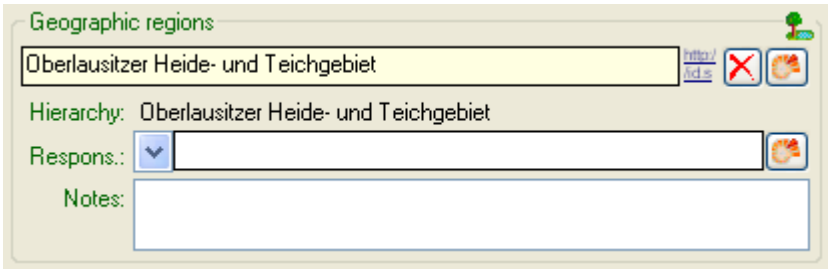
For the description of the habitat select the collection event or an existing collection site property . The description is entered in the field **Description of the habitat** (see below).



To enter a new property of the collection site, select the collection event or an existing collection site property and then choose the type of the property you want to enter from the dropdown menu as shown below. Only items that are not already set for a collection event will appear in the list. You can customize the selection of visible items.



The list is dependent on the availability of terminologies for site descriptions. Details for the property can be entered if you choose the item in the hierarchy. The person responsible for the entry as well as notes can be entered in the corresponding fields as shown below.



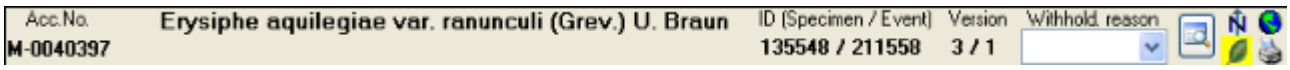
To delete entries use the button. You can enter a value by either typing the name of the collection site property or by selection from the module DiversityScientificTerms. To search for properties from this module click on the button. As responsible user the name of the current user will be inserted. You may change this by either typing or selection from the module DiversityAgents.

The data concerning the habitats are stored in the table CollectionEventProperty.

Specimens

Specimens are the entities stored in a collection. For handling the data use the appropriate buttons (new specimen, copy, delete). See the data section for further details. Directly attached to the specimen are the data about accession, label, exsiccatae, notes, reference and the availability. A specimen may be composed of several units. Any problems concerning the specimen should be entered into the problems field.

The header of the main window shows some important parameters for the specimen selected. On the left side the **accession number** is shown. In the center you find the last identification of the main identification unit according to the display order. The next field shows the internal **ID**'s (field CollectionSpecimenID in table CollectionSpecimen and CollectionEventID in table CollectionEvent) and the **Version**s for the specimen and the collection event. For details about the version of a dataset see the version topic. The availability of a specimen can be changed by entering an appropriate reason in the field **Withholding reason**.



If the specimen is a type, the header will show the type state and the identification connected with the type (see below).

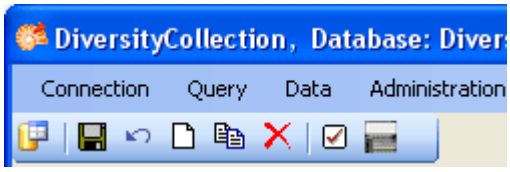
To inspect the history of a specimen click on the button. For further details see the history section.

With the buttons you can control the upper part of the window with the maps, the images for the collection event and the specimen as well as the label print. The buttons with the corresponding sector visible in the window are depicted with a red background. If there are images available and they are hidden, the background will turn to yellow.

The data are stored in the table CollectionSpecimen.

Data

The controls for the handling of datasets in the database are located in the left upper part of the window as shown in the image below. A step-by-step introduction for the creation of new datasets is provided in the tutorial.

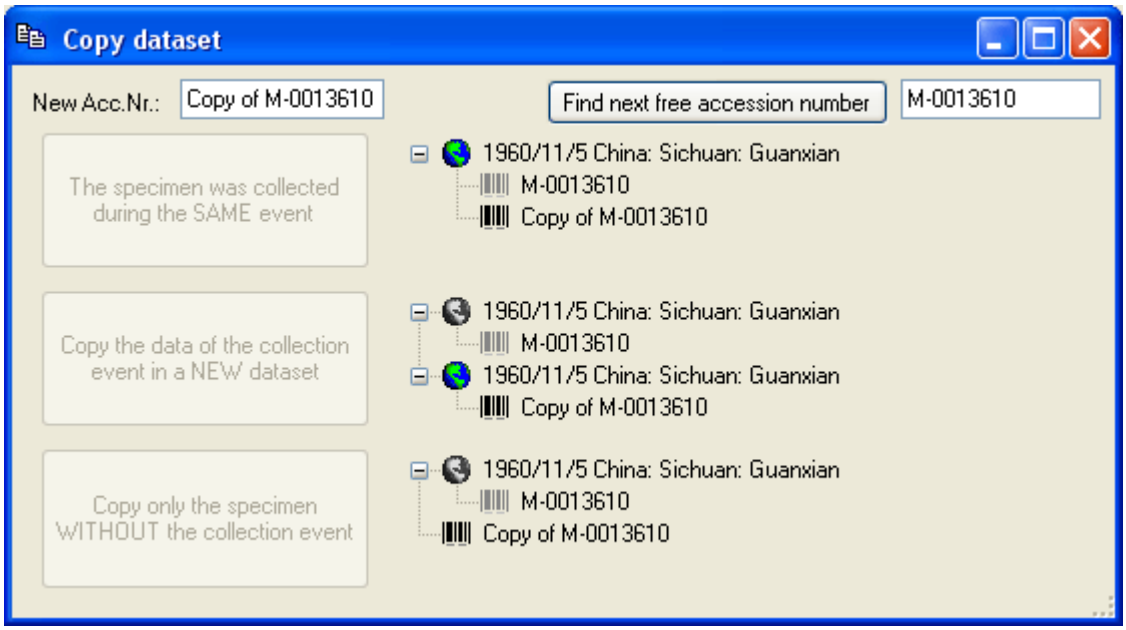


To **save** the changes in a dataset click on the button. If you select another dataset from the result list, the current changes will be saved automatically.

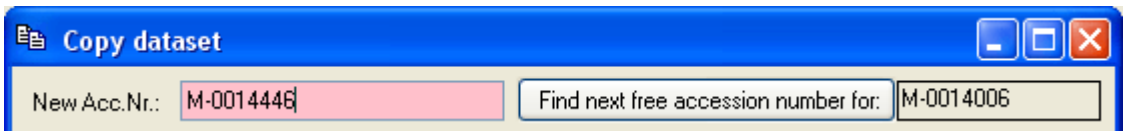
To **undo** the changes in a dataset click on the button. This will recover the original data unless the changes had been saved or changes were done in certain tables or hierarchies were the data must be stored to display the hierarchy.

To create a **new** entry in the database, click on the button below the search result listbox. This will create a new record of a specimen and show it in the result list.

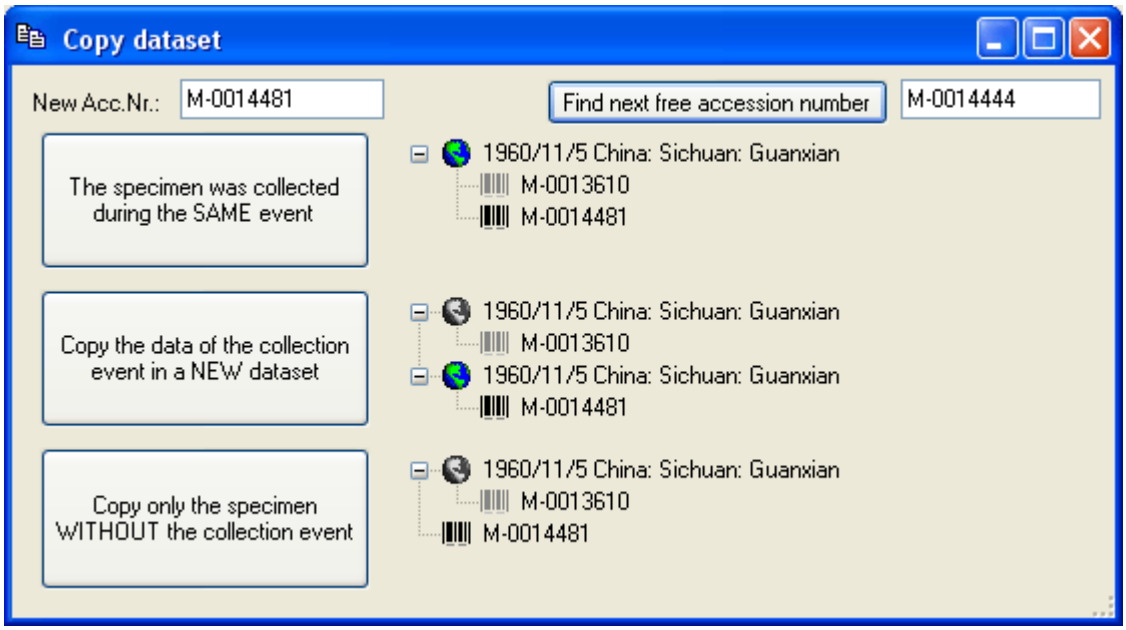
To **copy** the data of a specimen record, choose it from the list and click on the button . If the specimen you want to copy is linked to a collection event, a window will open and provide you with several possibilities for the copy as shown below.



The field for the new accession number contains a string like Copy of ... . If you change this the accession number to a value that is already present in the database you will be warned as shown below.



Use the **[Find next free accession number for:]** button to search for the next free number for the accession number shown in the second field. If the program finds a free number, it will be used as shown below.

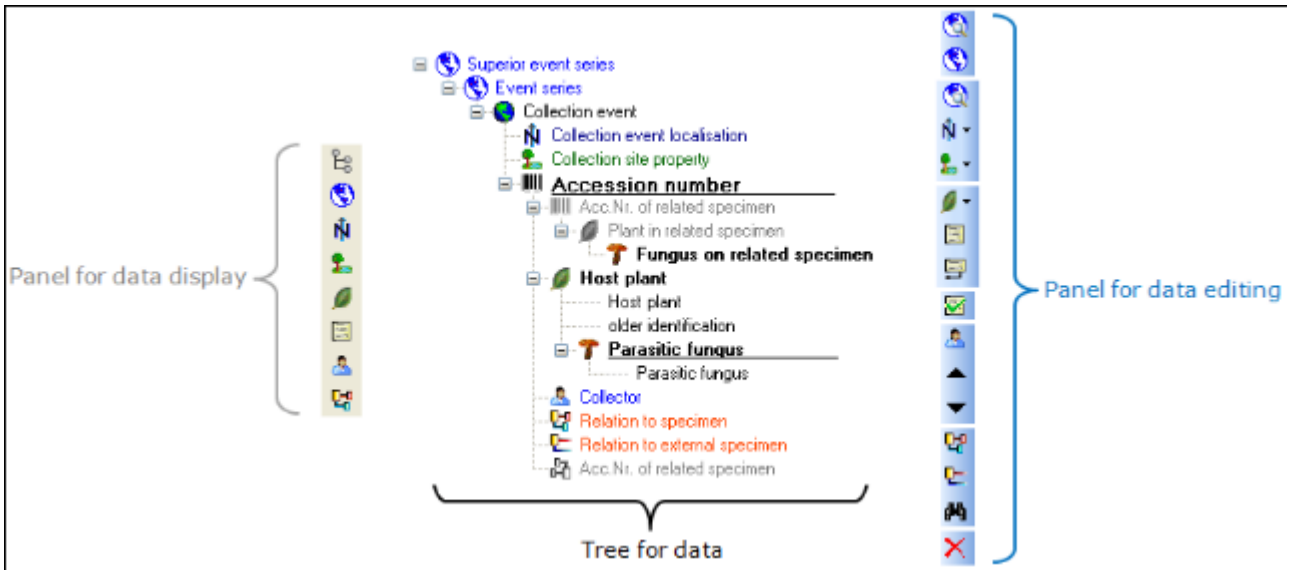


The first option will take the new dataset in the same collection event as the original data, so no new data for a collection event will be generated. The second option will create a copy of the original data for the collection event and connect the copy of the specimen record with the new event. The last option will only copy the specimen data and establish no connection to any collection event. If you decide not to copy anything, just close the window.

To **delete** a dataset click on the button ✕.

Tree for the specimen

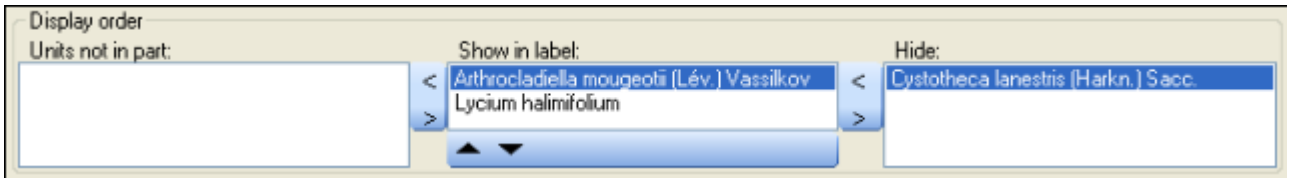
The upper tree in the window provides an overview for all the data linked with the specimen. With the buttons in the left panel you can hide or show certain nodes in the tree. So if for example you do not want to see the collectors, just click on the button and they will be hidden in the tree. The button will change to a grey version and the background will turn to yellow to show that there are hidden data of the collectors. The first two buttons (,) are visible if the collection event is part of a collection event series. They switch between two display modes for the event series. If you click on the button only the superior event series will be shown. To show the whole hierarchy of the event series, click on the button.



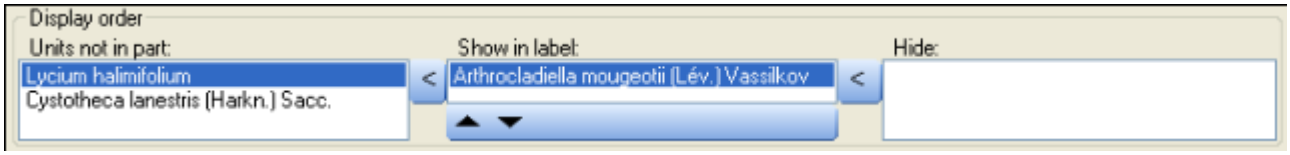
The panel on the right side of the tree is for editing the data, like for example the button will insert a new collector. For further details see the special sections.

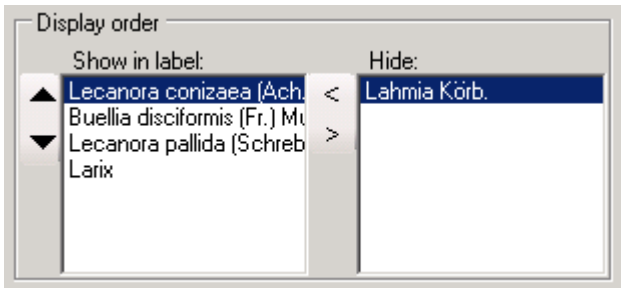
Display order

The display order defines the sequence in which the units within this specimen will appear on e.g. a label. The first unit will be printed in the header of the label, others are included in the text below. You can change the display order by using the ▲ ▼ buttons. If a unit should not appear on the label, transfer it to the hide list using the > button. The < button returns it to the list that will be shown on a label. The first unit can not be transferred to the hide list.



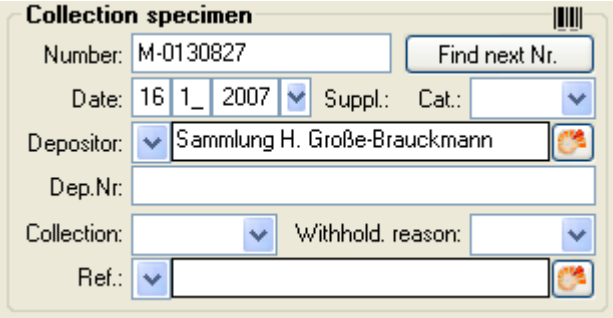
In addition to the display order you can specify whether an organsim is present in a selected part of a specimen.





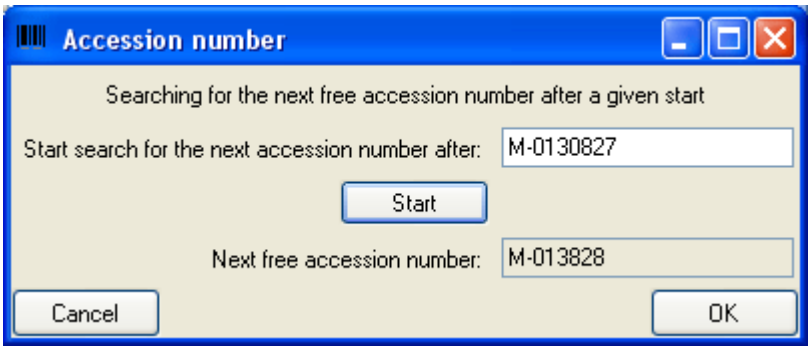
Accession

The accession of a specimen in a collection is documented with its accession number and if available the date of the accession.



If the specimen was received from another collection, this may be documented with the name of the depositor and the number in the original collection (Dep.Nr).

If a new specimen is entered you can use the Find next button to search for the next free accession number. A window will open as shown below, where you can start the search. The search will start with the number you give as a starting point.

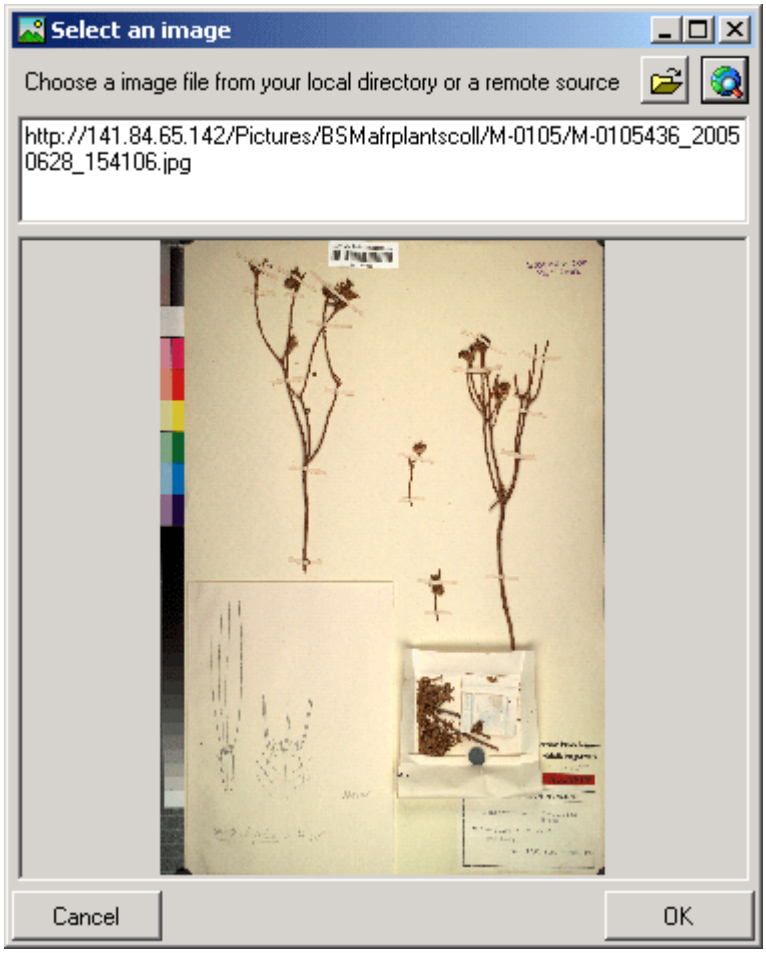


The system will try to find the next free number on the basis of the accession numbers available in the database. Click OK to use the number for the specimen.

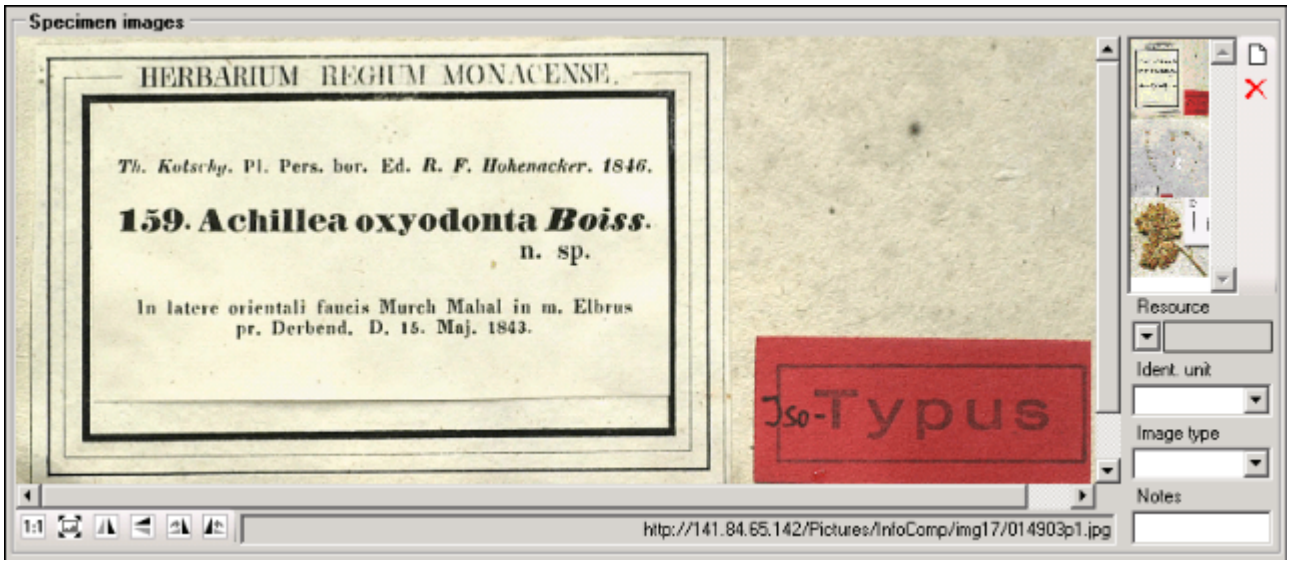
Data concerning the accession and deposition are stored in the table CollectionSpecimen.

Specimen images

To see the specimen images activate the icon in the image selector . If you choose the option **View - Show existing images** from the menu, the images will automatically be shown. Each specimen may be documented with several images. The images may be stored local with its path or as a reference to the module DiversityResources within the Diversity Workbench. To enter a new image, click on the button. A window will open where you can enter the path and file name of the image. Click on the button to search in your local directories or on the button to search for a web address. The selected image will be shown in the preview.



To delete an image, select it from the list and click on the button.

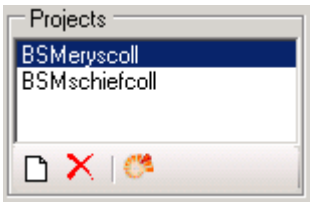


To zoom a sector of the image, just drag the mouse over the image. A red square will indicate the zoomed area. To set the size of the image to the original resolution click on the 1:1 button. To adapt the size of the image to the available space in the form click on the button. To change the orientation of the image use the appropriate buttons (flip horizontal, flip vertical, rotate right, rotate left). To view the image in a separate form, click on the button. If the image should not be published e.g. on the internet, enter any reason in the **Withhold. reason** - field.

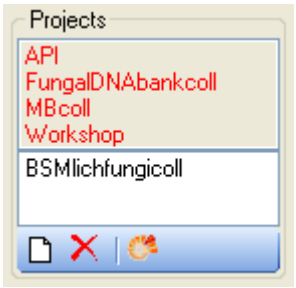
Data are stored in the table CollectionSpecimenImage.

Projects

Every collection specimen can be assigned to any number of projects. To assign a specimen to a project click on the button. To remove it from a project, select the project from the list and click on the button.

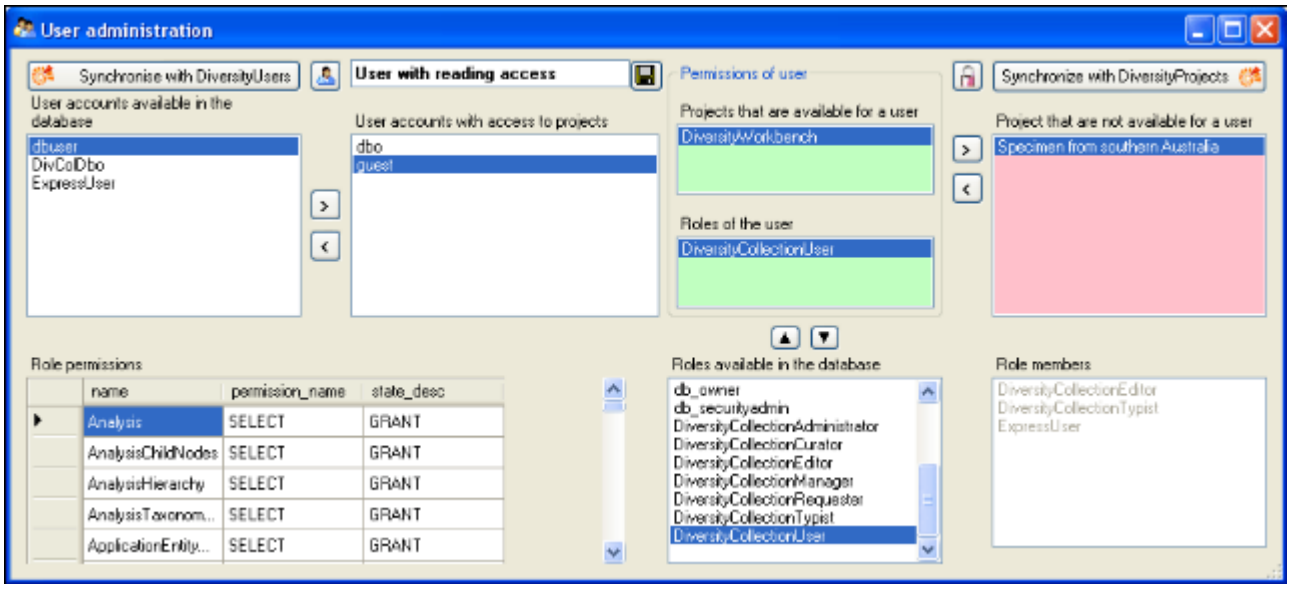


If there are projects, to which you have no access to, these will be listed in a separate list at the top as shown below.



Data are stored in the table CollectionProject.

Details upon the projects within the Diversity Workbench are stored in the database DiversityProjects. To open a project to see further information upon a project click on the button. To edit details in the projects you need the application **DiversityProjects.exe** in your application directory and access to the database DiversityProjects. To synchronize the projects listed in DiversityProjects you can use the synchronize functionality in the user administration window as shown below. If DiversityProjects is not available you can create a new project with the button. If DiversityProjects is available, use the synchronize functionality Synchronize with DiversityProjects.



Notes and problems

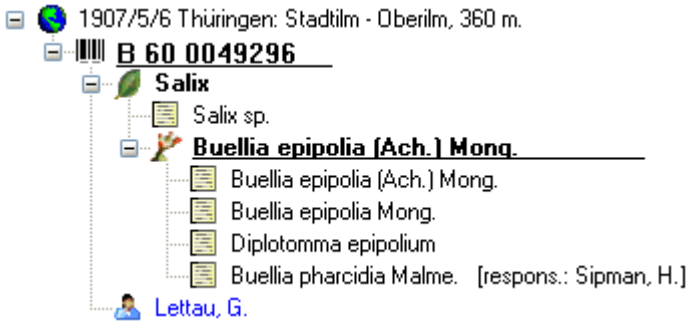
To enter notes or problems connected with the specimen select it in the hierarchy. The data form will then open the corresponding fields. The **Original** notes are the notes found on the label of the specimen, made by the original collector or from a later revision. **Additional** notes are those made by the editor of the specimen record, e. g. doubtful identification or locality.

In the **Problems** area enter the description of a problem that occurred during data editing. Typically these entries should be deleted after help has been obtained. Do not enter scientific problems here. Use Additional notes for such permanent problems!

The data are stored in the table CollectionSpecimen.

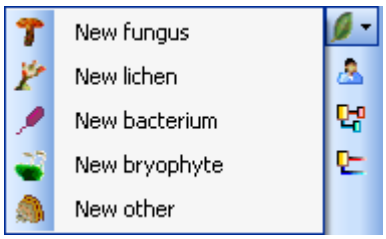
Organisms and identifications

Each specimen can contain several organisms and each organism may have been identified several times as shown in the image below.



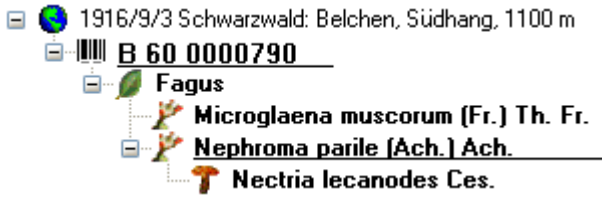
Identification unit

The items or organisms in one collection specimen are regarded as identification units. One specimen can contain several identification units, e.g. an insect (1) feeding on a fungus (2) growing as a parasit on a plant (3). To add a new organism use the drop down menu as shown below to select the taxonomic group to which the new organism belongs to.

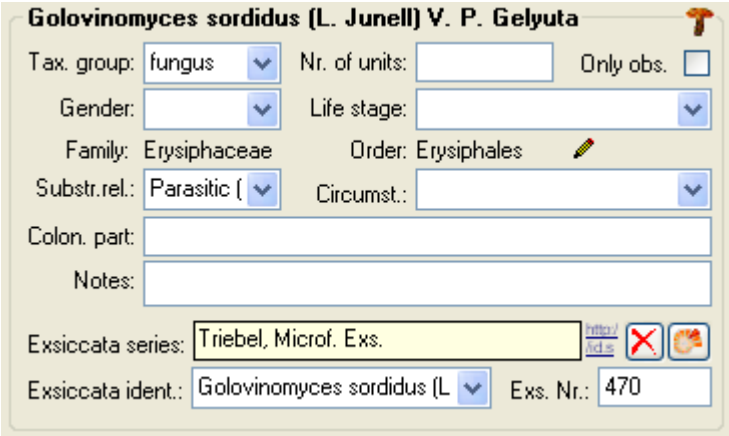


To specify the taxonmic groups that are shown in the drop down menu select Administration - Customize display ... from the menu.

The organisms of the specimen are shown in the tree. To edit the relations between the organisms just do this by drag and drop. The name of the organism under which the specimen is stored in the collection is underlined. To delete an organism select it in the tree and click on the ✕ button.



To enter details about one of the organisms like the e.g. the gender or the life stage, select it in the tree. Then the fields for the details of this organism are shown in the right area of the window as shown below.



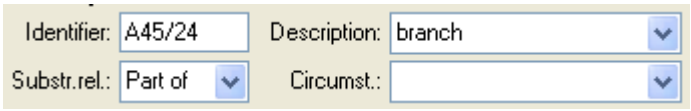
Taxonomic hierarchy - family and order

The entries for the family and the order of the taxon are either set when linking to a taxonomic database or manually when no link to a taxonomic database exists.

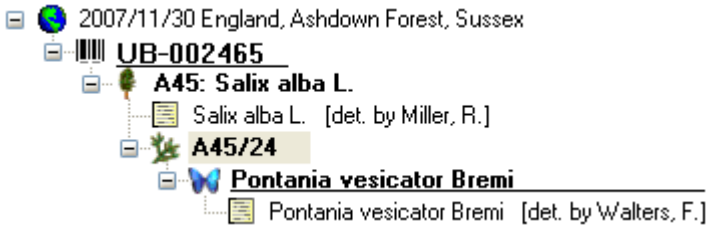
If no link to a taxonomic database exists you may enter the family and the order after clicking on the ✎ button. To transfer these entries to other specimens with the same genus use the maintenance functions as described under Maintenance - family and order.

Parts of units

If a unit contains parts that have to be documented, you can do this by setting the relation to the substrate to "Part of".



In the tree the backgroud of a part will be grey as shown below for the unit-tree und the tree depicting the storage of a sample. The name of the part will correspond to the identifier set for this part and the icon will correspond to the description of the part provided you choose one of the preset options contained in the drop down list. You may of course enter any description for the unit. A unit that is part of another unit can not get an identification. Identifications are restricted to the main unit.

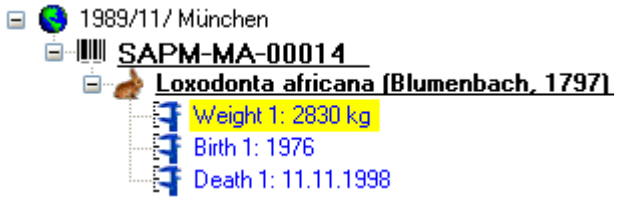


For details about exsiccatal series and analysis see the related topics. You can sort your identification units e.g. for display on a label with the display order. Each identification unit can have several identifications.

The data for the organisms are stored in the table IdentificationUnit.

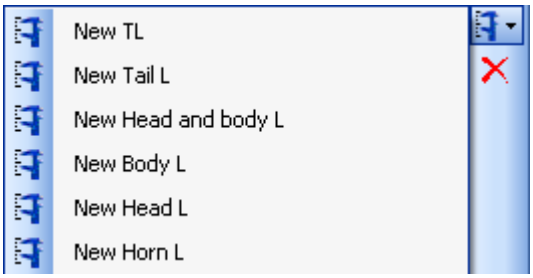
Analysis

The organisms in a specimen can be analysed by analysis types defined in the database. In the tree the analysis entries are symbolized with an icon as shown below. Only the types of analysis that were assigned to the group of the organism can be selected. An analysis always referes to an organism and may refer to a part of a specimen. An analysis that refers to a part of a specimen will be shown in the tree for the parts as shown in the examples below.

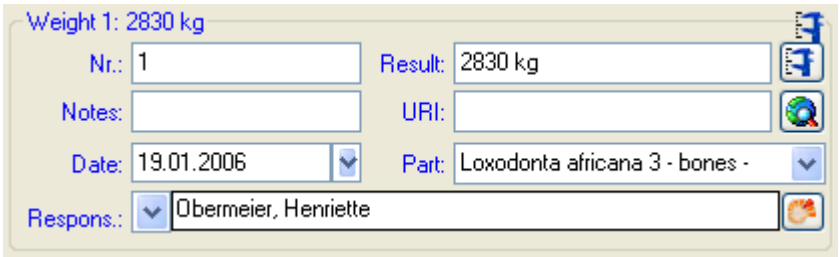


To show or hide the analysis items in the tree use the button in the left panel. This button has 3 states. In the default state the analyis entries will be shown as above. If you click a second time on the button it will change to the hierarchy mode and the entries will be shown as in the tree below. The last state will hide the entries.

To insert a new analysis select the organism which has been analysed in the upper tree. Then select the type of the analysis from the drop down list as shown below.



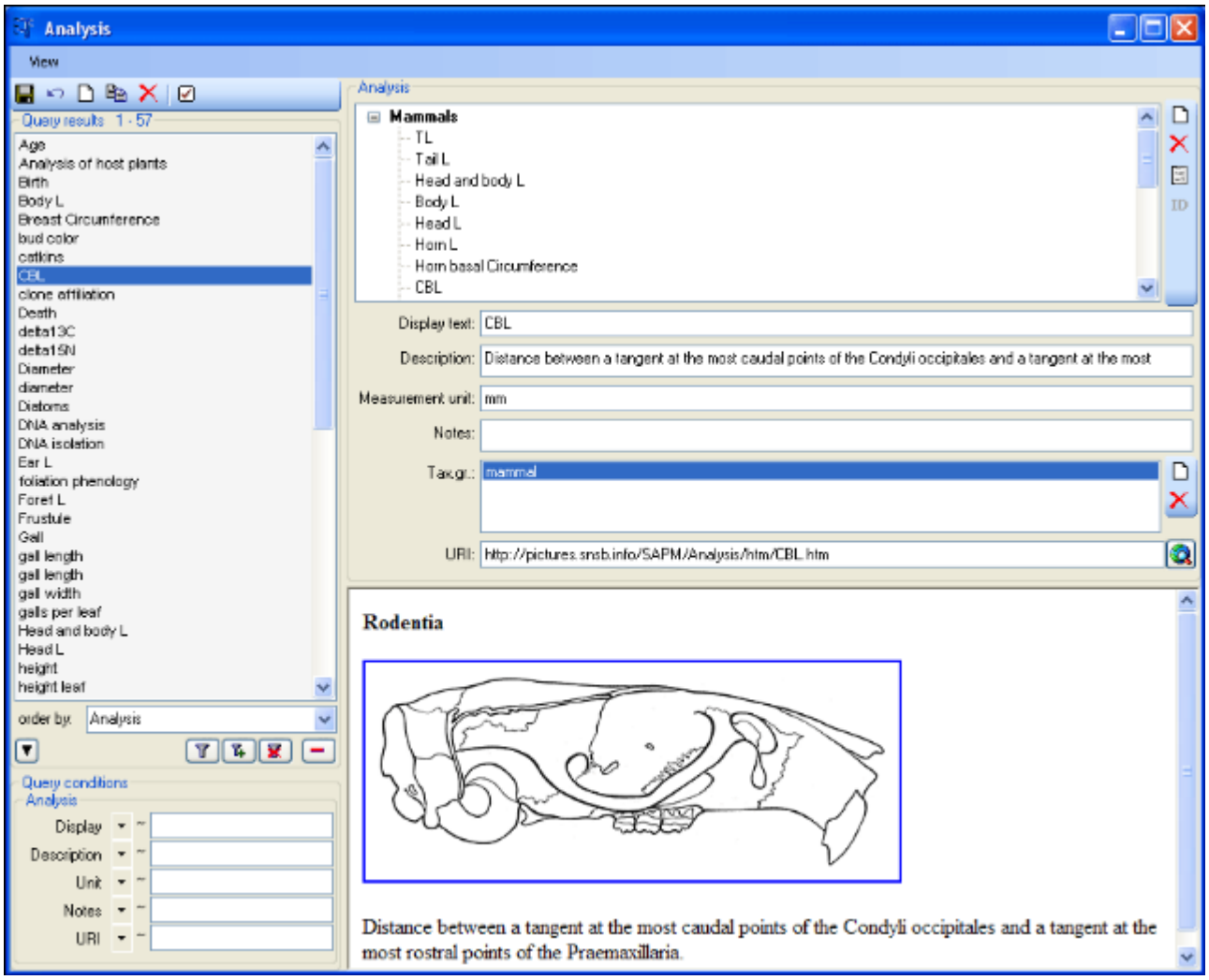
To delete an analysis, select it in the tree and use the button in the panel on the left. To edit the details of an analysis, select it in the tree to enter the fields as shown below. To enter or inspect an URI given for a single analysis, click on the button.



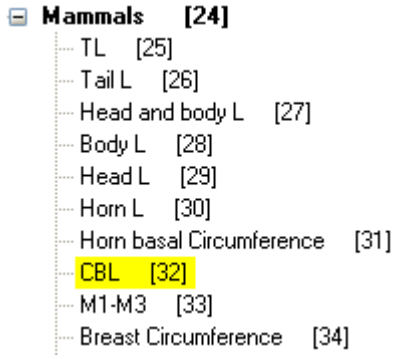
If an analysis refers to a part of the specimen, you can document this by choosing the

respective part from the list (see field **Part** above). The analysis will then be shown in the tree for the parts of the specimen as well. Alternatively you may directly choose the organism in the part tree and create the analysis for this part of the organism in the part tree.

If you need further information about an analysis click on the button to open the window for the analysis administration. If you have the proper rights you can edit the analysis types used in your collection as shown below. This window is also accessible via the menu entry **Administration - Analysis...**



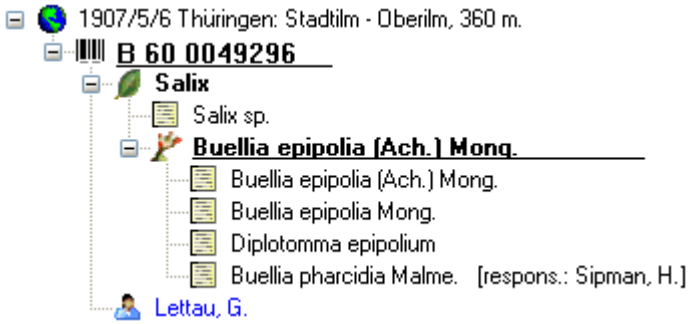
For the import and export of data it is sometimes necessary to know the ID's of the analysis types. To see the ID's, click on the **ID** button. Than the ID's will be shown as in the image below.



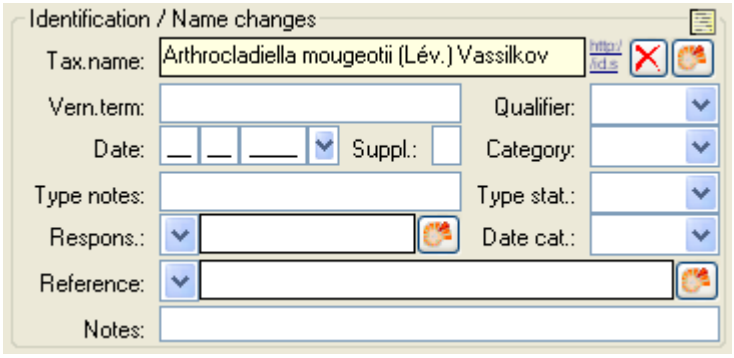
The types of an analysis are restricted by the taxonomic group of the organism or object to be analysed. Edit the list of taxonomic groups that can use a certain analysis using the and button. For details about handling the data see the data section. If you want to or inspect an URI given for an analysis type, click on the button. In the window below the URL related to the analysis can be shown.

Identification

Each organism or unit may have been identified several times. The identifications are listed in the tree under the organism as shown in the image below. To create a new identification choose the organism from the tree and click on the icon. The last identification will always be taken as the valid one and set at the top of the list. To insert an older identification at the base of the list click on the icon.



To hide / show the identifications in the tree click on the icon on the right panel next to the tree. To enter a confirmation of a identification, choose it in the tree and then click on the button. To delete an identification select it in the tree and click the button. To enter details for an identification, choose it in the tree. You then can enter the details in the form opening on the right side of the tree as shown below.

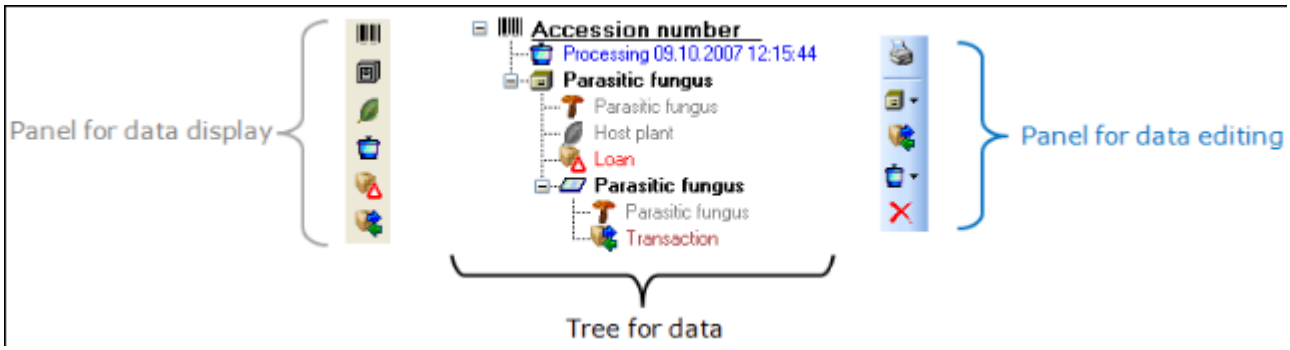


The taxonomic name of an organism can be selected from either the names already available in DiversityCollection or from the module DiversityTaxonNames. To choose from the local names type the beginning of the name and click on the drop down button to get a list of the available names. You can use wildcards according to SQL.

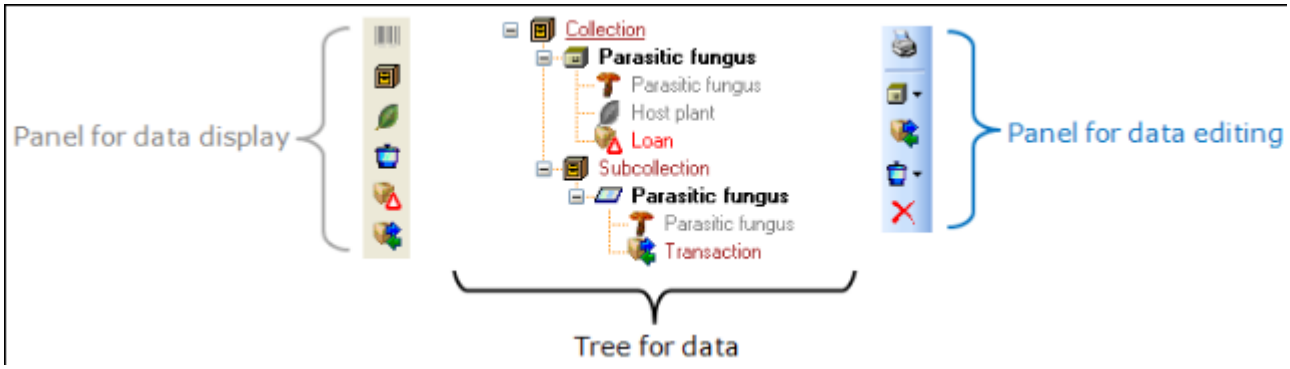
To select a name from an external database for taxonomic names, click on the button. A window will open where you can search for a taxonomic name. In the database the data for the identification are stored in table Identification.

Storage and trees for the parts of the specimen

In the tree at the base of the window the parts of the specimen with their related data are shown. This tree depicts the parts either in accordance to their relation to each other in the specimen tree (click on the icon in the panel on the left) ...



... or depending on the collections where the parts are stored (click on the icon in the panel on the left).

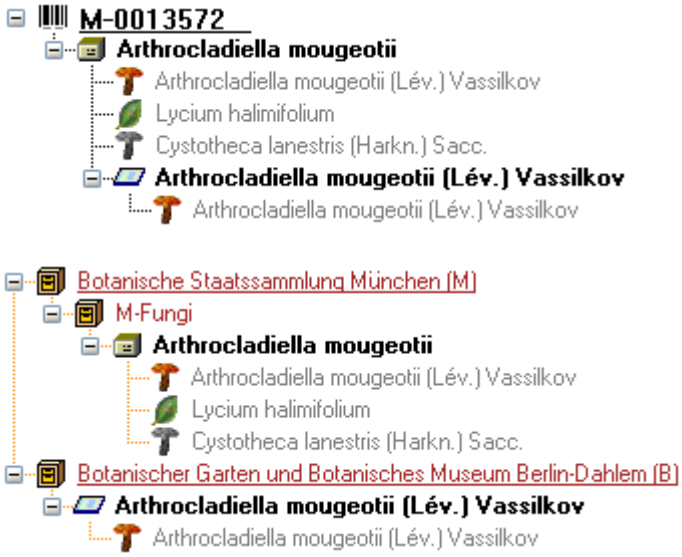


With the buttons in the left panel you can hide or show certain nodes in the tree. So if for example you do not want to see the processings, just click on the button and they will be hidden in the tree. The button will change to a grey version and the background will turn to yellow, to show you that there are hidden data of the processings.

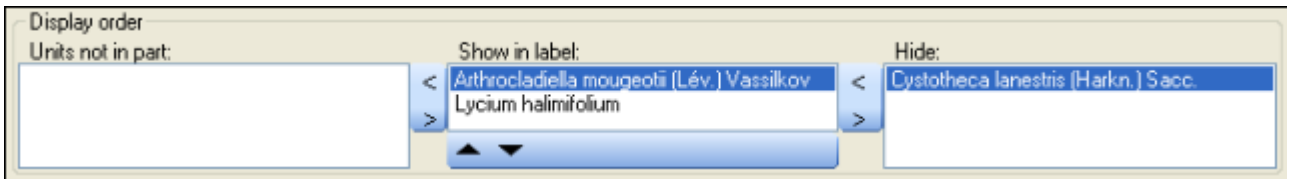
The panel on the right of the tree is for editing the data, like for example the button will insert a new transaction. For further details see the sections Transaction and Processing .

Parts of the specimen

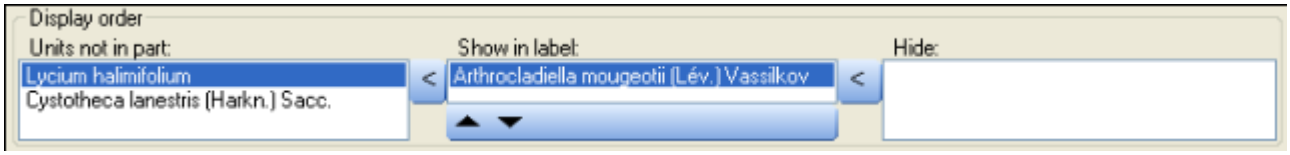
A specimen can contain several parts which may be stored in several collections. These parts may represent different material categories, e.g. herbarium sheets, microscopic slides etc. The parts are shown in the part tree under the main tree. This tree depicts the parts either in accordance to their relation to each other in the specimen tree (click on the icon in the panel on the left) - see first image below - or in relation to the collections where the parts are stored (click on the icon in the panel on the left) - see second image below where the two parts of the specimen are located in two collections. To add a new part, use the drop down field in the panel on the right of the tree and choose the material category of the new part. The material categories that appear in this list can be customized. You can set the default collection for new parts. If the default collection is not set, you will be asked to name the collection of the new part. If the part is derived from another part, select this in the tree before creating the new part. The new part will then be a child of the old part. You can later edit these relations by drag & drop. If you want to create a copy of a part, use the button in the panel right of the tree.



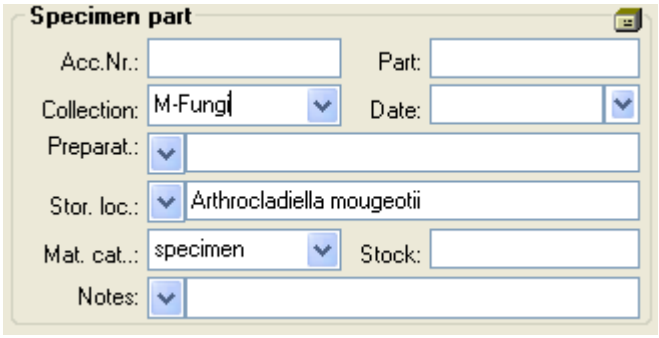
If an organim is present in a part of the specimen, it will be shown in the tree. To edit the presence, select the specimen part in the tree and edit the **Display order** as shown below.



In the expample above, all organisms are present in the specimen, but the fungus *Cystotheca lanestris* was chosen to not appear on the label and transferred to the **Hide** list. This is symbolized by a grey icon in the tree. The slide, symbolized by the icon, was created from the sample, so in the specimen tree it is shown as a child of the specimen symbolized by the icon. Concerning the organisms, only the fungus *Arthrocladiella* is found on the slide. So in the tree it is the only organ ism shown as child of the slide. In the display order the two other organisms appear in the **Units not in part** list (see below).



To move the organisms between the lists use the **>** and **<** buttons. In the list **Show in label** you can change the sequence of the organisms with the ▲and ▼buttons in the panel at the base of the list. The organism at the top will be the one that will be taken as the main organism, e.g. for printing a label. To edit the details of a part, select it in the tree to open the fields as shown below.



The identifier, e.g. the taxonomic name under which the specimen can be found in the collection is entered in the field **Storage location**. You can search for this entry in the query.

To enter the **storage location** you can use the drop down button to select from the identifications within the specimen. The drop down button for the Notes will provide you with a list of all entries in this field, filtered with your entry in the text field. You may use wildcards like "%" or "_" (see wildcards).

In the collection tree, the specimen are to shown with their dependence upon each other, but where they are located with the collections. To get informations about a collection just select it in the tree. If you have the necessary rights, you can use the button to open for the administration of the collections. This form is as well accessible under the menu entry **Administration - Collections...**

The material category of a specimen part is indicated by the icon in the hierarchy.

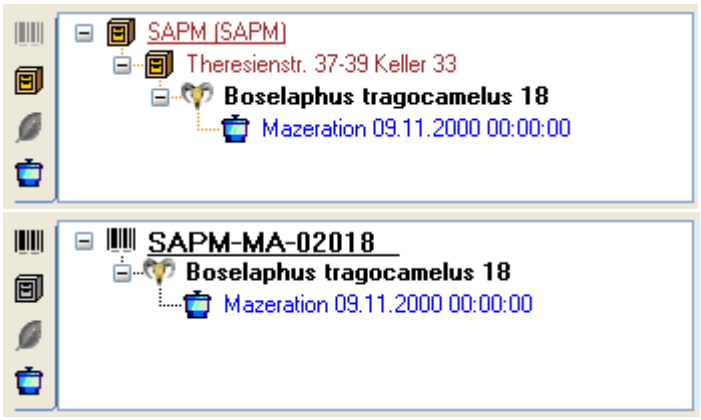
Here some expamples:
bones: bones or skeleton from vertebrates
cultures: living cultures of organisms
drawing: original line or color drawing
herbarium sheets: capsules or sheets as stored in a botanical collection
icones: icones, images etc. stored in a botanical collection
micr. slide: glass plate with sections of specimen for microscopic studies
specimen: specimen stored in a collection

Specimens can be included in a transaction and every stored part of a specimen can be treated with several processings.

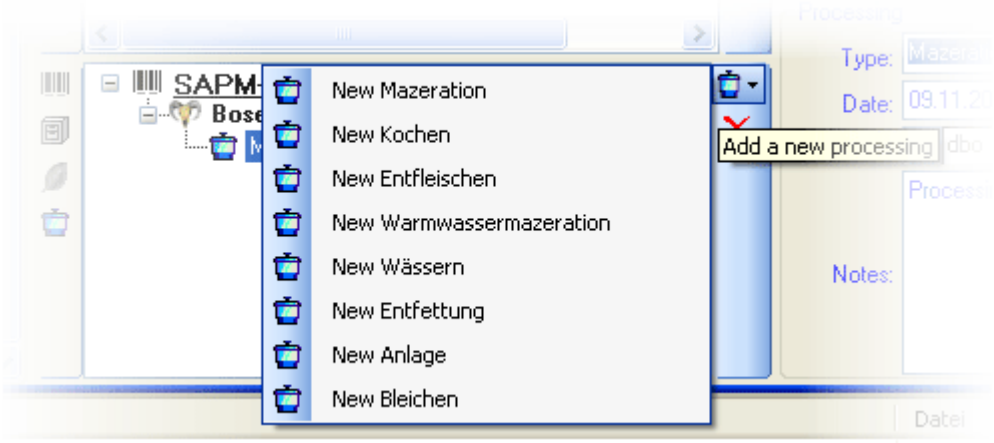
Data are stored in the table CollectionSpecimenPart.

Processing

Every specimen can be processed, e.g. for preparation or preservation. Processings can be applied to a specimen or a part of a specimen. In the hierarchy for the parts the processings are indicated by the icon and a blue text.If there are any processings this will be indicated by the icon in the tool bar. You can hide or show the processings in the hierarchy with a click on the icon. The images below show a processing in either the collection or the specimen tree for the parts.

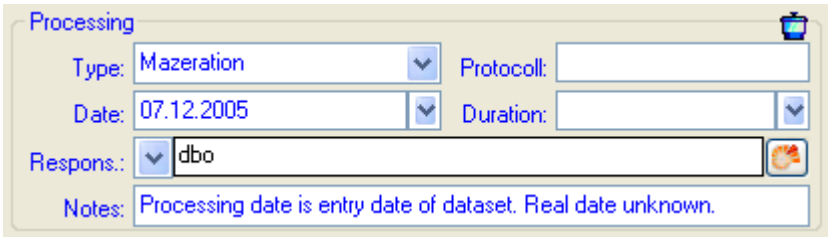


You can enter a new processing for either the whole specimen or a part of the specimen. To do this select the specimen or a part in the hierarchy and then choose a processing from the list as shown below.

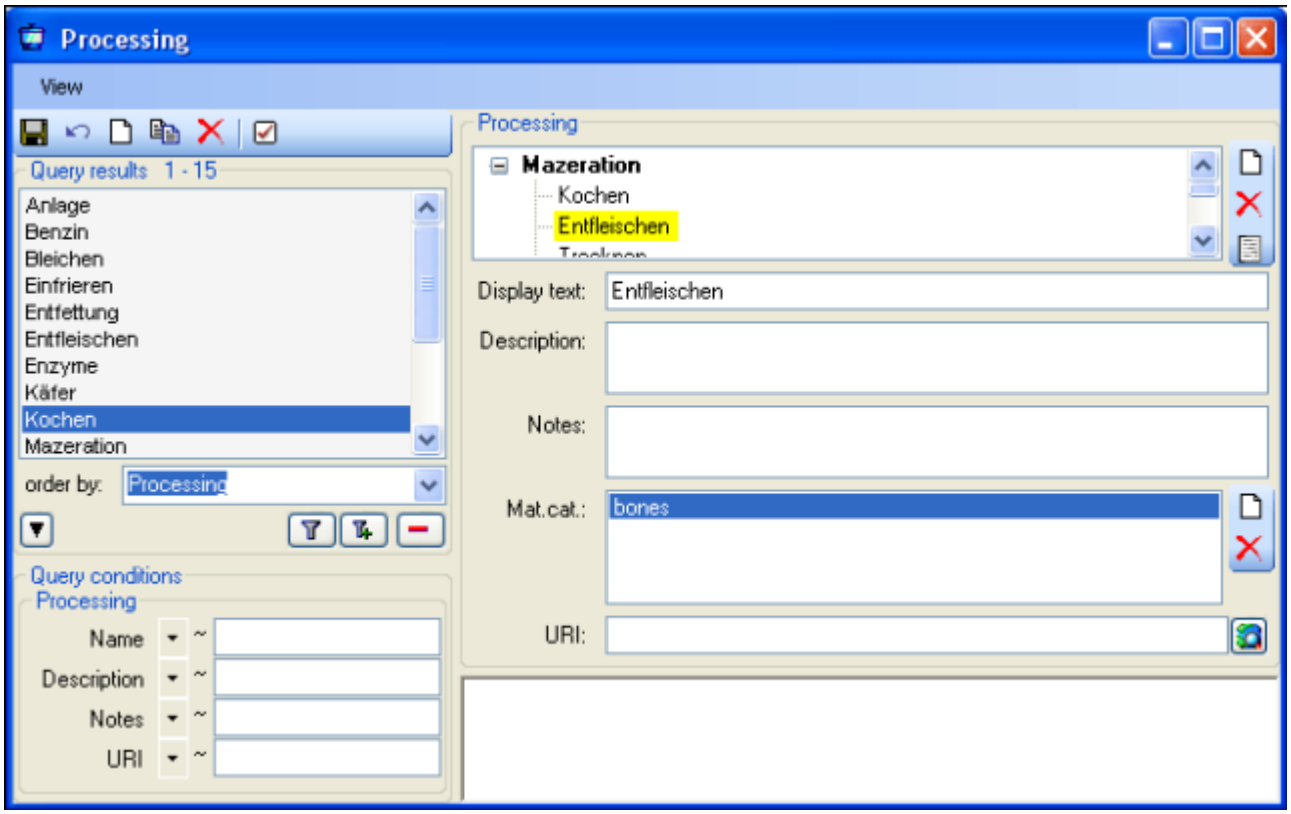


To delete a processing select it and click on the button. To edit the data for a processing select it in the hierarchy. The window will show you the fields related to the processing as shown below.

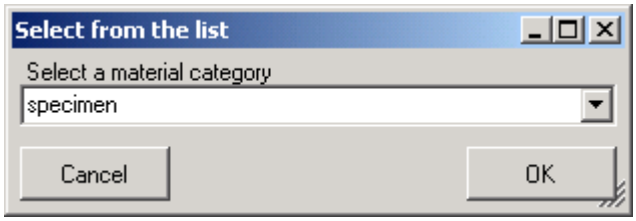
In the window select the type of your processing from the list.



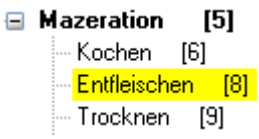
If you have the proper rights you can edit the processing types under the menu entry **Administration - Processing...**



In this window you can define the processings used in your collection. To add a new processing click on the button, to delete a processing select it and click on the button. The processings are related to material categories within the database. To add a new material category that should be connected to the selected processing, click on the button in the panel right of the material categories. A window will open and you have to select the material category that should be connected with the processing (see below).



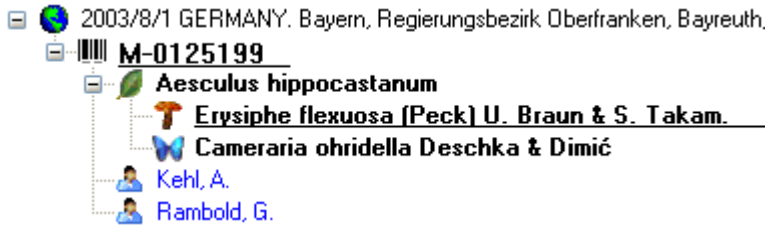
For the import and export of data it is sometimes necessary to know the ID's of the analyis types. To see the ID's of the datasets, click on the **ID** button. Than the ID's will be shown as in the image below.



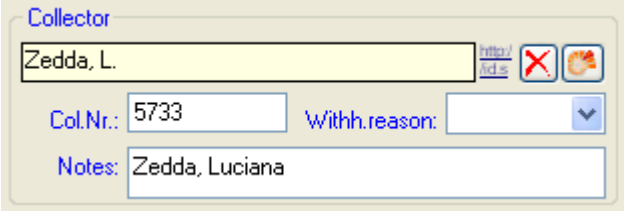
The data of the processing are stored in the table CollectionSpecimenProcessing.

Collector

The people or groups responsible for the gathering of the specimen are stored in the table CollectionAgent. There may be several collectors for one specimen. In the tree the collectors are symbolized by the icon as shown below.



To hide or show the collectors in the tree use the icon in the left panel (it will change to a grey version with a yellow background when the authors are hidden). To insert a collector, choose either the specimen or an existing collector in the tree and then click on the icon in the right panel. To enter details for a collector, select it in the tree. In the right part of the window the fields for the details will then displayed (see below).



If a collector should refer to a dataset in the module DiversityAgents (where more details like addresses may be stored) click on the button. The sequence of the collectors (e.g. for print on a label) as shown in the tree can changed by with the ▲ and ▼ buttons in the panel on the right of the tree.

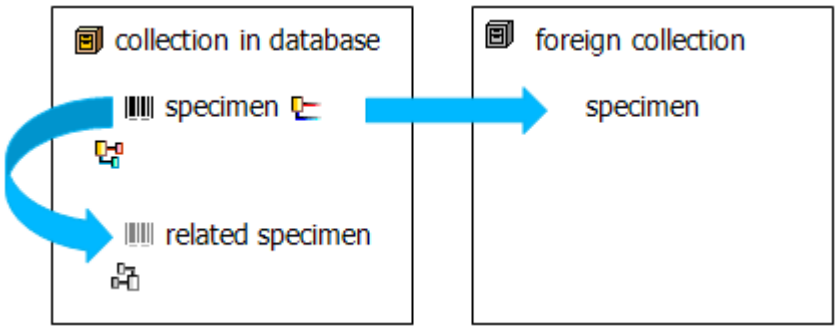
To remove a collector, select the entry in the tree and click on the button.

You can search for the gathering numbers of the collectors in the query.

The data of the collectors are stored in the table CollectionAgent.

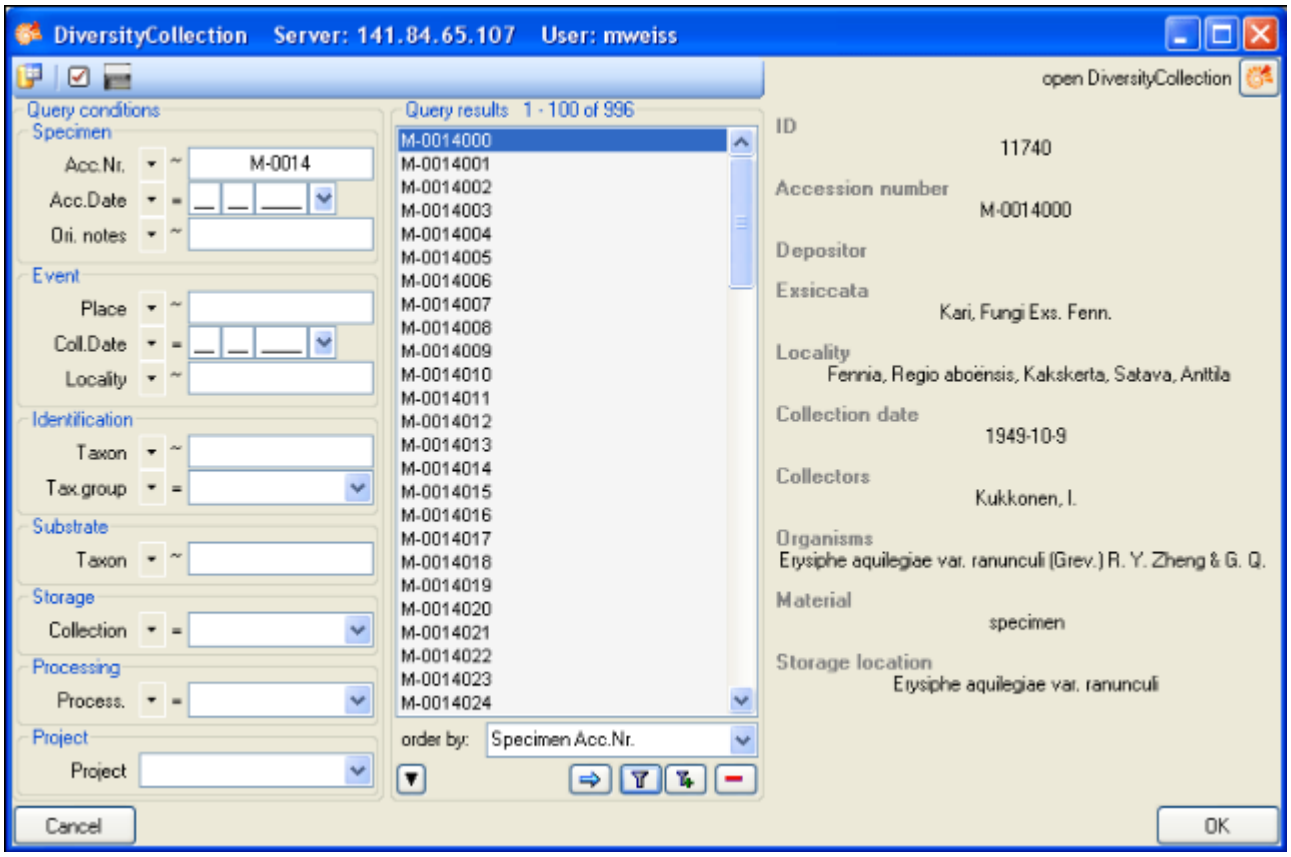
Relation

Each specimen may have relations to other specimens. This can be documented by adding relations to these specimens. To add a relation select the specimen in the tree (symbolized by the barcode). Then you can either add a relation to a local specimen using the button or a relation to a remote specimen using the button. To show or hide the relations in the tree use the and buttons.

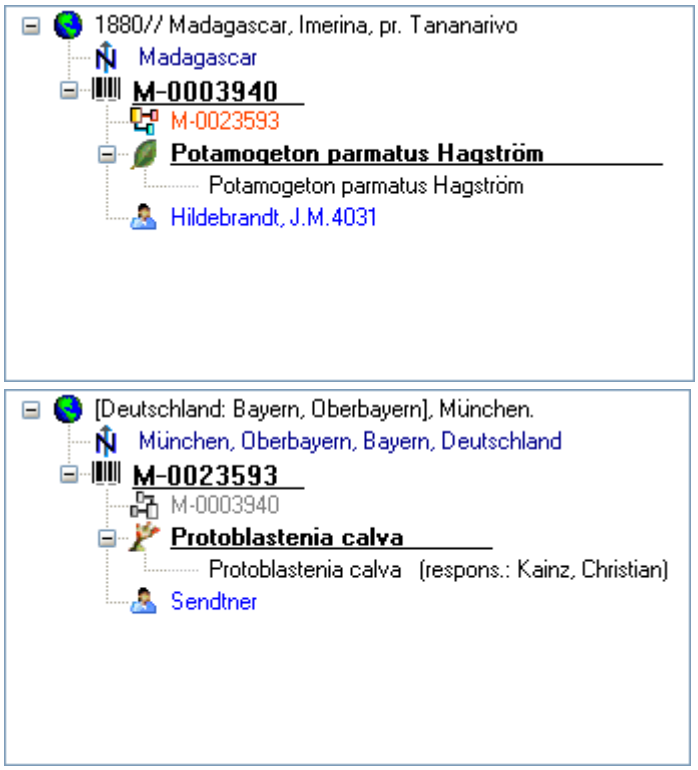


Internal relation

To add an internal relation to a local specimen use the button. A window will open where you can search for the related specimen.



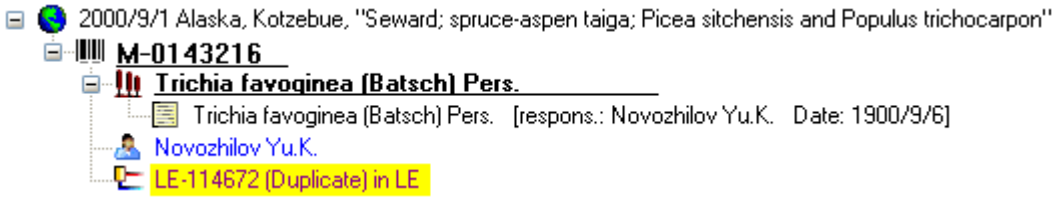
Select the related specimen from the query result list and click OK to close the window and establish the relation. In the left image below you can see an internal relation , while the image on the right shows a reverse relation from another specimen.



In the case of internal relations you can navigate to the related specimens using the button.

External relation

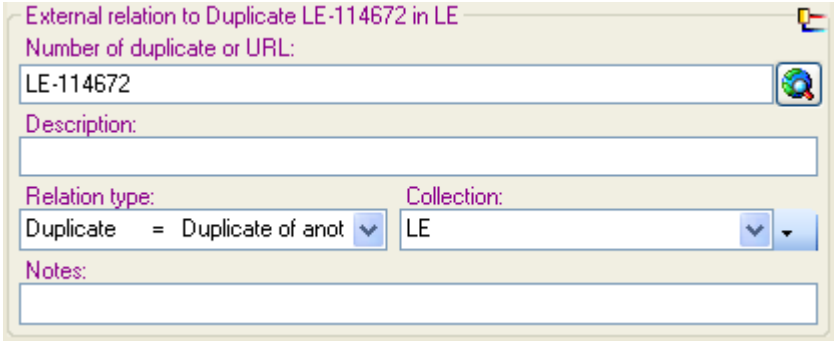
To add an external relation to a specimen that is not administrated in the database DiversityCollection use the button. In the image below you can see a relation to a specimen in a external collection.



To delete a relation select it in the tree and click on the button.

Editing the data

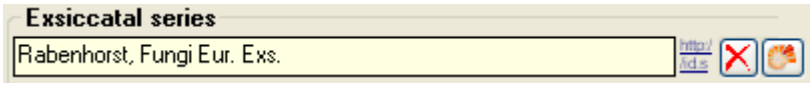
To edit the data for a relation, click on the entry in the tree. In the right part of the window the details of the dataset will be shown (see below).



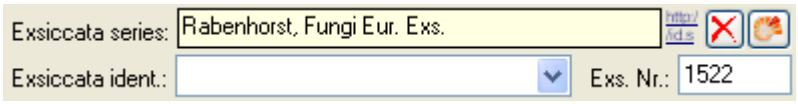
The data are stored in the table CollectionSpecimenRelation.

Exsiccatae

A specimen may be part of an exsiccatal series. These series are administrated in the Diversity Workbench module DiversityExsiccatae. To enter the exsiccatal series select the specimen or the organism resp. identification unit in the tree. With the specimen the exsiccatal series is shown at the base of the data form as shown below.



As there may be several organisms with separate exsiccatal numbers in one specimen the numbers are handled together with the identification units. Select the concerned organism to enter data in the form as shown below.



You can either type the name of the exsiccatal series or choose one from the module DiversityExsiccatae. If you click on the A window will open showing the link to the entry as derived from the module.

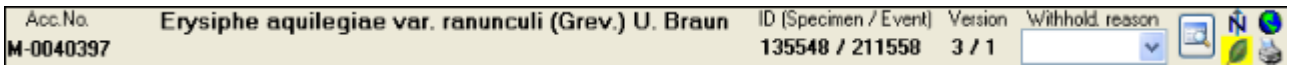
The data about the exsiccatal series are stored in the table CollectionSpecimen, the data about the number and the taxon are stored in the table IdentificationUnit.

Access to the data

To get access to the data, you have to take several hurdles. In DiversityCollection, you must be a member of one of the user groups. You have only access to those data, that are listed in the projects you have access to. For external users, data may be blocked by entries in the data witholding reasons.

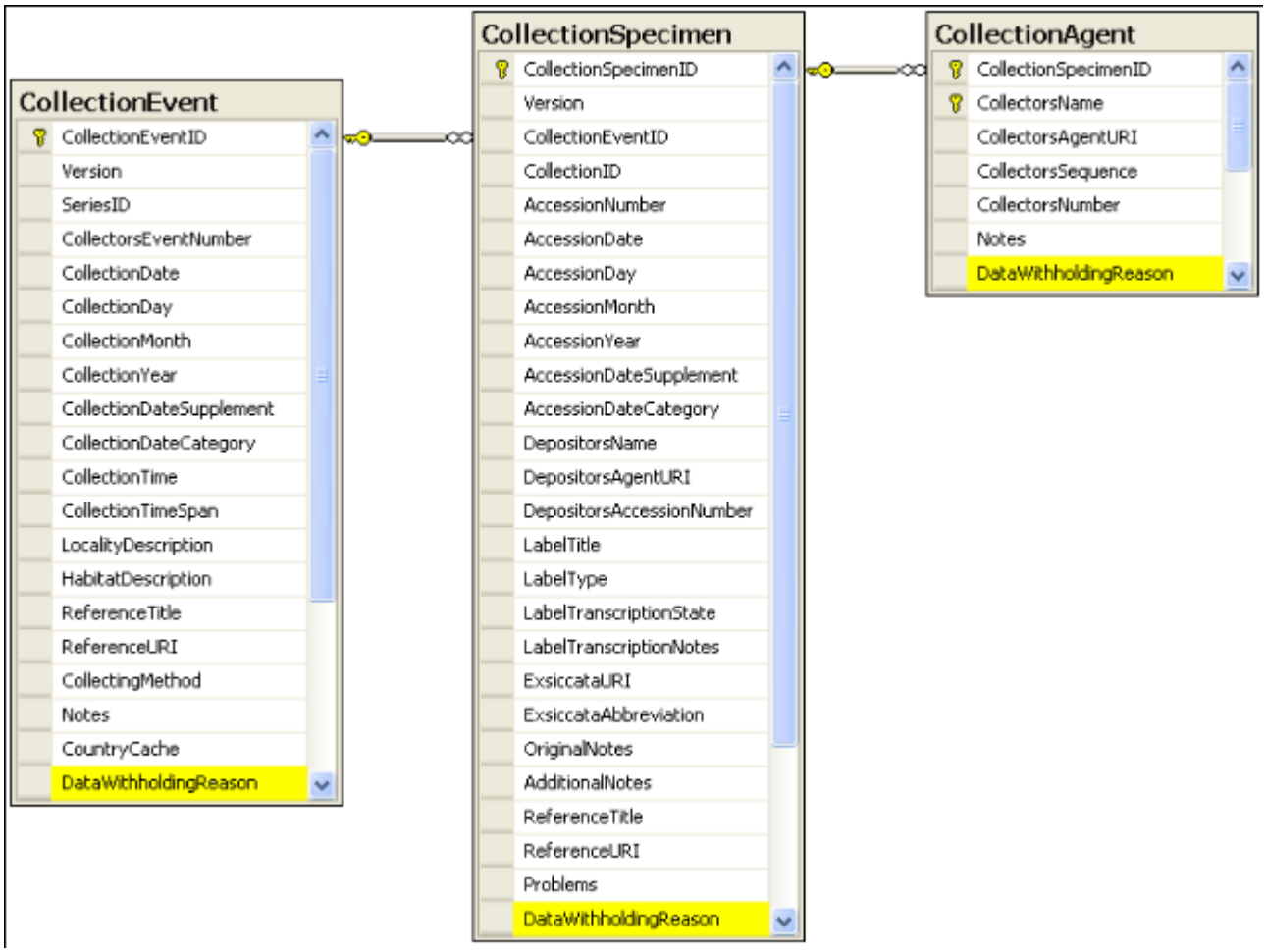
Availability of datasets

The data of certain parts of the database can be blocked for remote access. If you want to prevent access for the whole dataset, enter your reason in the field **Withholding reason** in the header (see below).



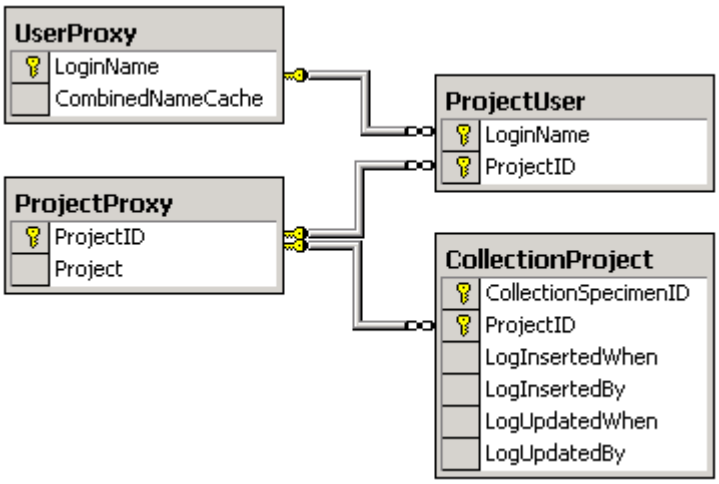
If you want to block only the access to the data of the collection event or the collector choose them in the tree and enter your reason for withholding the data in the corresponding fields. The data will only be published if these fields are empty.

Within the database a user has only access to the data of those projects for which he has the permission. If a dataset belongs to a certain project, this is stored in the table CollectionProject. To prevent a publication of the data these can be blocked for the whole dataset (table CollectionSpecimen), for the locality and collection site informations (table CollectionEvent) and for the collectors (table CollectionAgent) - see overview below - and for all image tables: CollectionSpecimenImage, CollectionEventImage, CollectionEventSeriesImage.



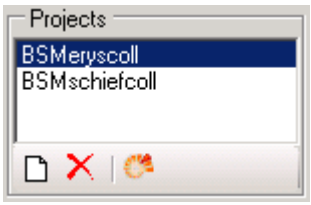
Permissions for projects

The access for the user to the data within the database are stored in the tables shown below. The tables UserProxy and ProjectProxy are related to the Diversity Workbench modules DiversityUsers and DiversityProjects respectively. See user administration for further details.

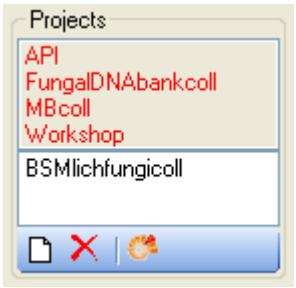

UserProxy
LoginName
CombinedNameCache
ProjectProxy
ProjectID
Project
ProjectUser
LoginName
ProjectID
CollectionProject
CollectionSpecimenID
ProjectID
LogInsertedWhen
LogInsertedBy
LogUpdatedWhen
LogUpdatedBy

Projects

Every collection specimen can be assigned to any number of projects. To assign a specimen to a project click on the button. To remove it from a project, select the project from the list and click on the button.

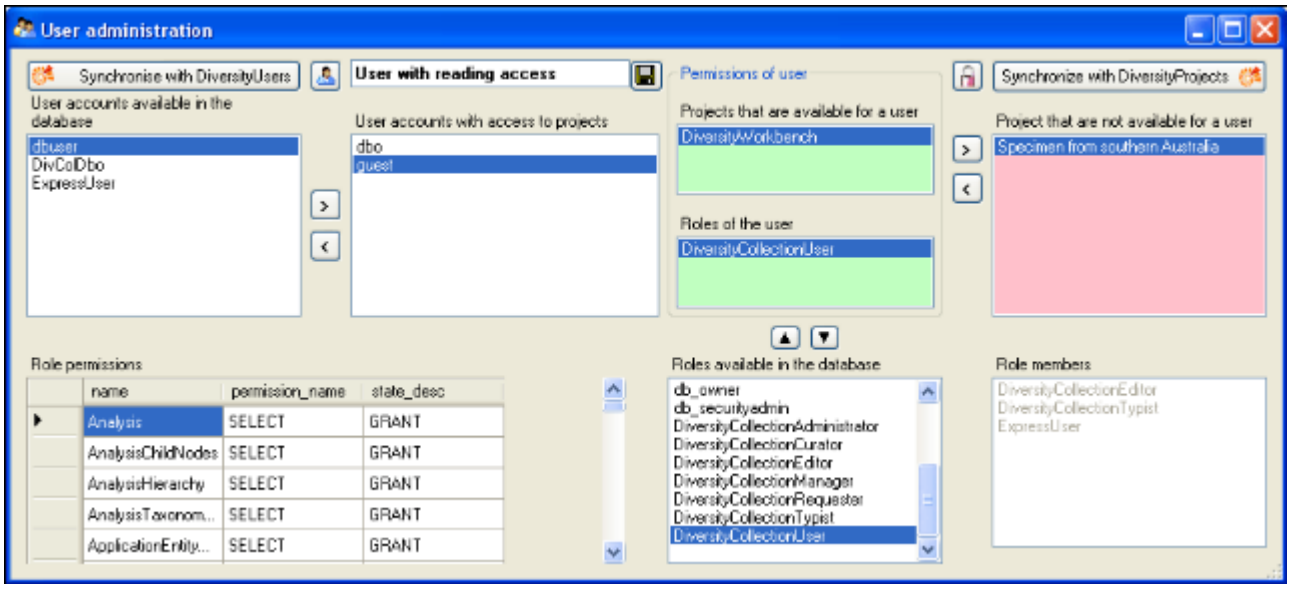


If there are projects, to which you have no access to, these will be listed in a separate list at the top as shown below.



Data are stored in the table CollectionProject.

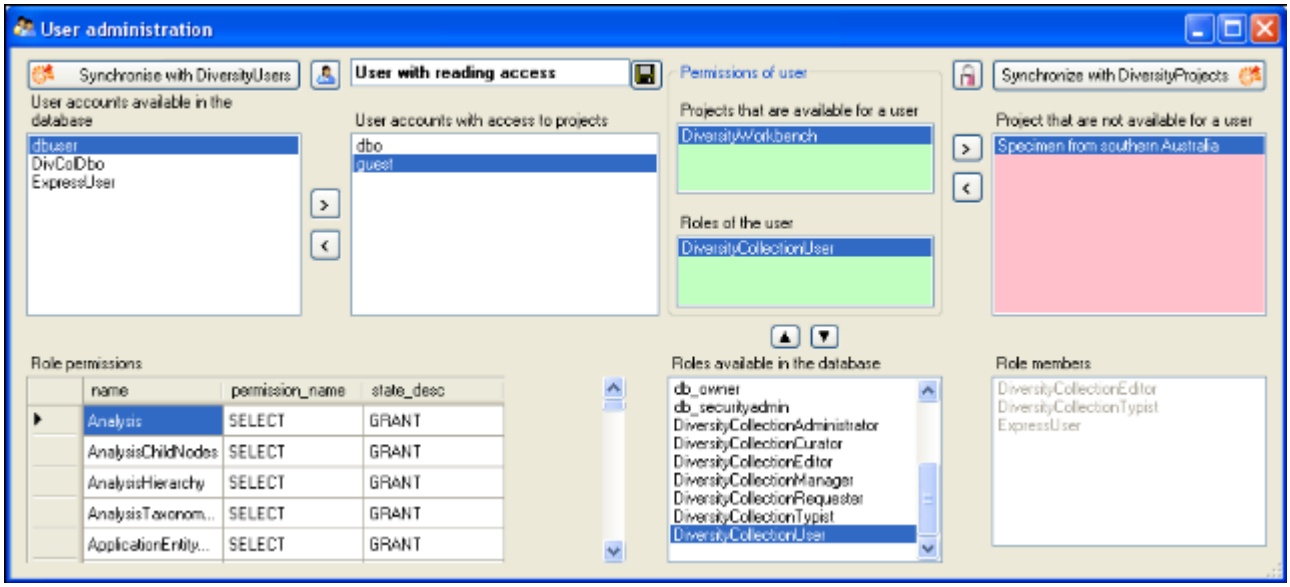
Details upon the projects within the Diversity Workbench are stored in the database DiversityProjects. To open a project to see further information upon a project click on the button. To edit details in the projects you need the application **DiversityProjects.exe** in your application directory and access to the database DiversityProjects. To synchronize the projects listed in DiversityProjects you can use the synchronize functionality in the user administration window as shown below. If DiversityProjects is not available you can create a new project with the button. If DiversityProjects is available, use the synchronize functionality Synchronize with DiversityProjects.



User administration

The permissions of users in the database are set via user groups resp. roles in the database and the access to the projects. To set the permissions choose **Administration** - **User** ... from the menu. A window as shown below will open. Here, among other administration tasks, you can change the permissions of the users. The first list shows the user accounts that are available in the database but have no access to any of the projects. To synchronize this list with the Diversity Workbench module DiversityUsers click on the **Synchronise with DiveristyUsers** button. To create a new SQL-Server user, click on the button. To create a new Windows user with access to the database resp. allow an existing user the access to the database use the Microsoft SQL Server Management Studio (see the installation section for further details). To permit access to a project click on the > button. To remove a user from this list use the < button.

The area **Permission of user** shows the projects a user has access to and the roles of the user within the database. To move users between the lists Projects that are available for a user and Projects that are not avialable for a user use the > and < buttons. To create a new project click on the button. If you use the Diversity Workbench module DiversityProjects, you can create a new project there and user the **Synchronise with DiversityProjects** button. To change the roles of a user use the ▲and ▼ buttons. Underneath the project list for a user you find the list of the Roles of the user. This list can be changed by using the and buttons. Underneath the roles list you find the list with the roles available in the database. On the left side of this list the permissions of the selected role are listed and on the right side the users with this role (Role members).



The permissions of the roles resp. user groups are shown in the lower left part of the form.

The role **DiversityCollectionUser** can see the data within the permitted projects, but can not change anything.

The role **DiversityCollectionTypist** has the same rights as the role DiversityCollectionUser and can edit a part of the user defined data.

The role **DiversityCollectionRequester** has the same rights as the role DiversityCollectionUser and in addition can place requests for specimens.

The role **DiversityCollectionManager** has the same rights as the role DiversityCollectionUser and in addition can handle transactions, i.e. shipments of specimen between collections and edit the collections for the collections he has the permission.

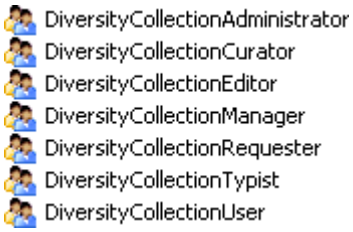
The role **DiversityCollectionEditor** can change the user defined parts of the data.

The role **DiversityCollectionCurator** has the combined rights of the roles DiversityCollectionEditor and DiversityCollectionManager.

The role **DiversityCollectionAdministrator** can delete data, edit the contents of internal tables, change user permissions etc.

Security

A user can be in 5 groups with diverse rights in the database where the higher groups have all rights of lower groups in addition to special rights for this group, e.g. DiversityCollectionUser can only read the data of certain tables while DiversityCollectionTypist has the rights of DiversityCollectionUser and additionally can edit the data in certain tables - see overview below.



Summarzied overview of the permissions of the groups

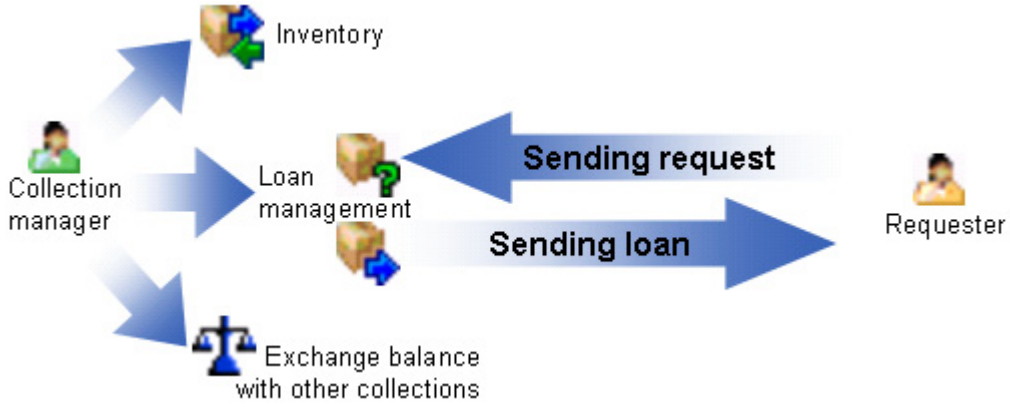
Role	**Permissions in addition to lower role resp. user group**
Administrator	Delete data, edit user permissions
Curator	Combines the roles editor and collection manager
Editor	Create new entries and delete details (not whole datasets)
Manager	Administration of collections, handling loans etc.
Requester	Has the right to place requests for specimen
Typist	Edit data
User	See the data of the data tables

If you are an Administrator you can add a user to one of these groups

Any user may have access to several projects.

Collection management

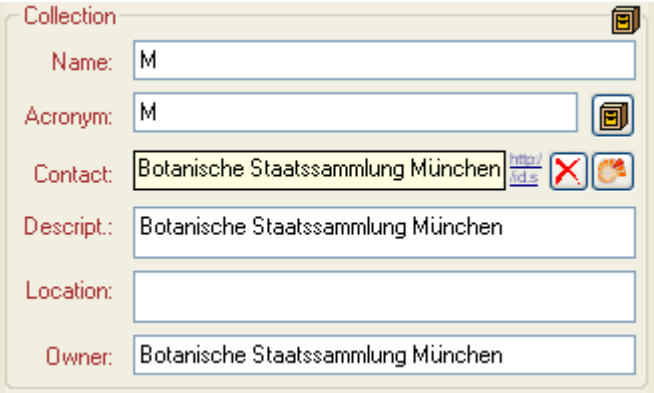
Collections are managed by **collection managers**, who organize the transfer of specimen between collections and track the **exchange balances**. Any transfer of specimen is organized via **transactions**. To be able to place are request for specimen from a collection, you must be in the group of **requesters** for this requested collection and a collection manager for the requesting collection. An overview is given in the image below.



Collection

The organisms in a specimen can be stored in several collections. To see the place where a specimen or parts of it are stored use the lower tree and click on the icon to select the display according to the collections as shown below.

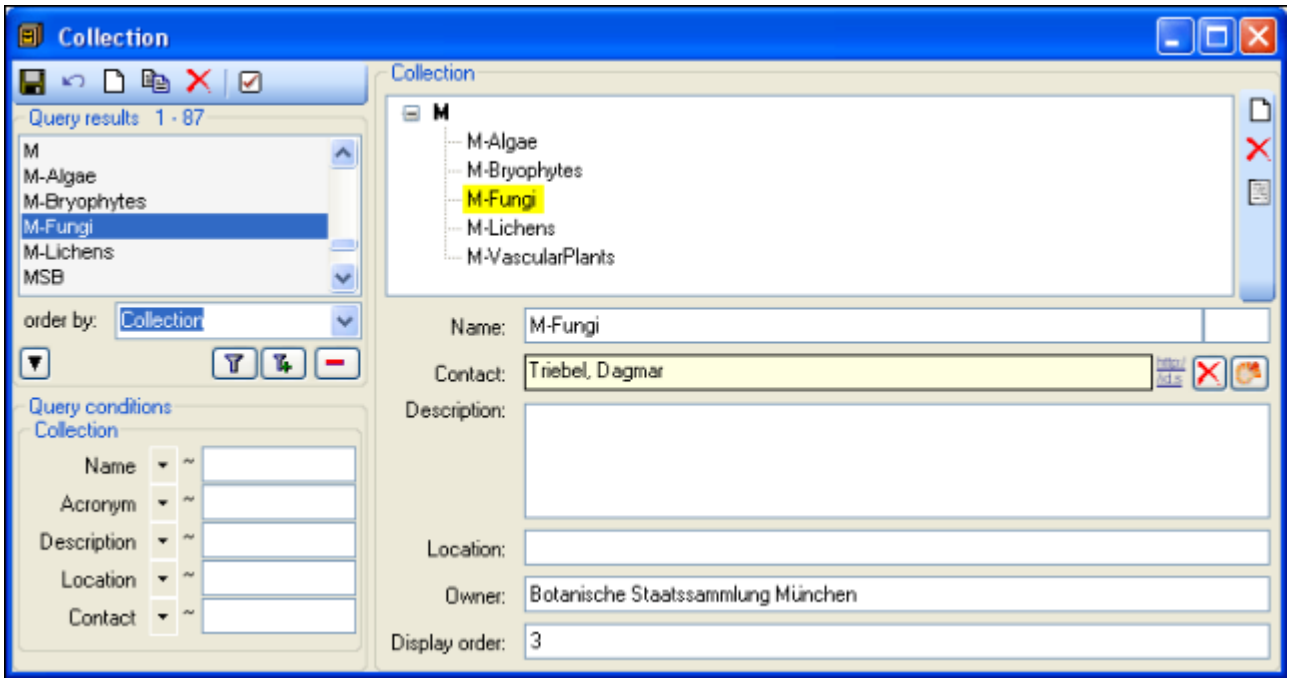
In the tree shown above select the collection to display the datafields as shown below.



To be able to edit the detais of a collection, you must be a collection manager of this collection. The restrictions for the access for editing the collections are defined in the collection manager part.

At the base of the fields you find the default collection, used for the creation of new specimen parts. Specimen parts that are not created as a part of another part will be placed in this collection. Use the button to edit this entry.

If you need further information about a collection click on the button or choose **Administration - Collections...** from the menu to open the window for the collection administration (this menu entry is only available for collection managers). This will open a window as shown below where you can edit the collections used in your database for which you are a collection manager.

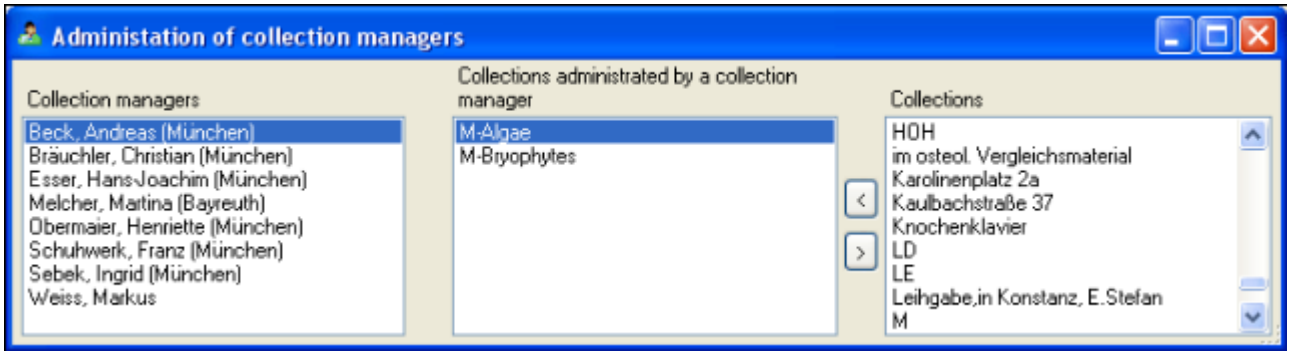


The specimens stored in a selected collection are listed in the specimen list. You can hide / show this list using the / button. To change to a specimen from this list click on the button.

For details about handling the data see the data section. Data of the collections are stored in the table Collection.

Collection manager

Collection managers are users with the right to edit the transactions. To edit the collection managers you must be in the administator group. Choose - **Administration** - **Collection managers** from the menu to open the editing window for the collection managers as shown below.

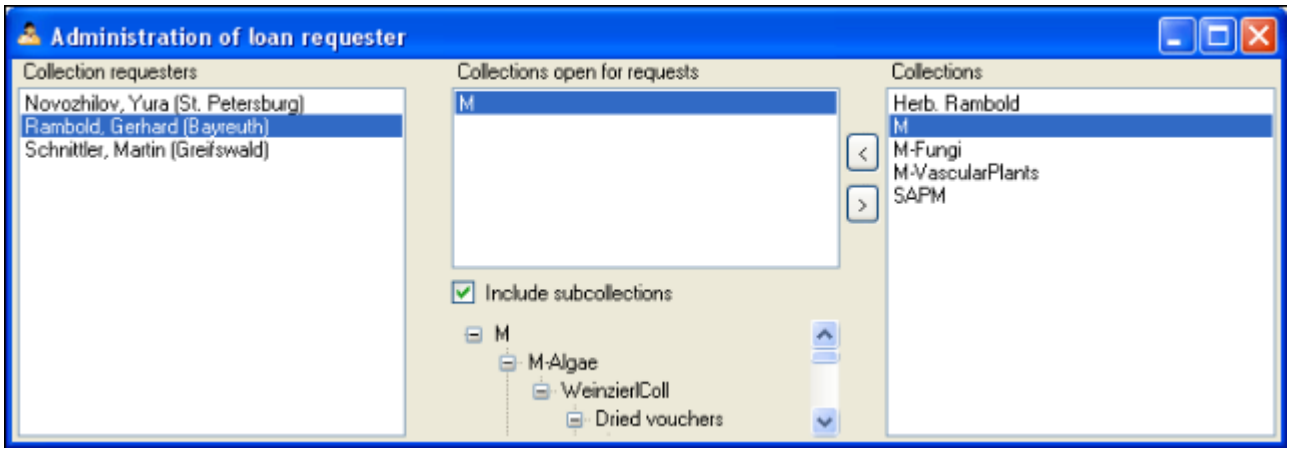


Use the **<** and **>** buttons to edit the list of the collections for which the selected collection manager can create and edit transactions.

For details about handling the collections see the transaction section. Data of the collections are stored in the table CollectionManager.

Collection requester

Collection requesters are users with the right to place requests for loans in other collections. To edit the collection requesters you must be a collection manager for the requested collection. Choose - **Administration** - **Collection requesters** from the menu to open the editing window for the collection requesters as shown below. The list on the left contains all users that are in the group Collection requesters. See user administration for further details. Select a user in this list to edit the accessible collections for this requester.

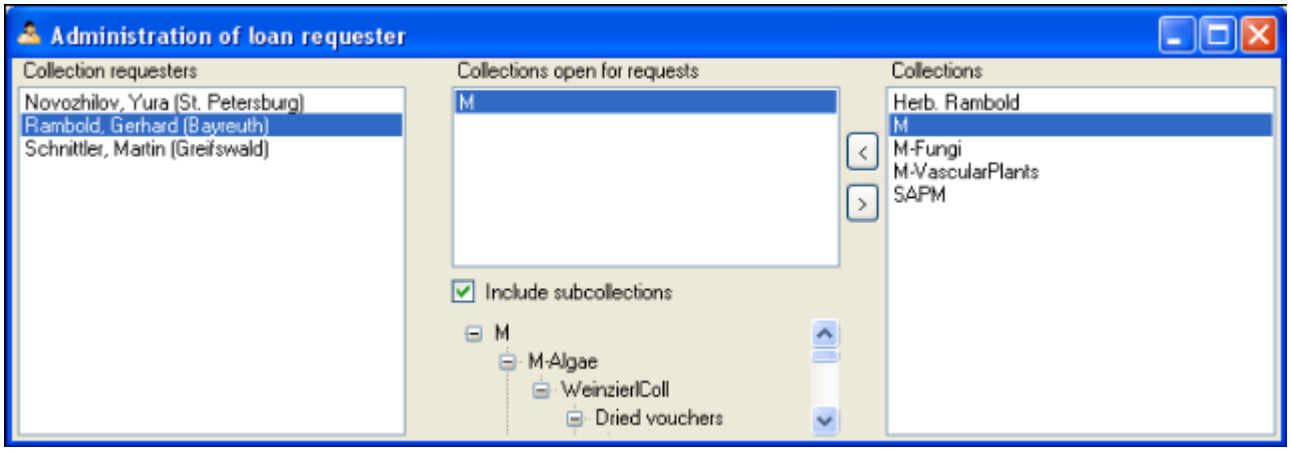


Use the **<** and **>** buttons to edit the list of the collections for which the selected requester can place a request. If the option Include subcollections is choosen, the requester can send requests for all subcollections of a collection. Otherwise the requests can only be sent for the collections in the list. The subcollections are shown in the hierarchy underneath the list.

For details about handling the collections see the transaction section. Data of the collections are stored in the table CollectionRequester.

Transactions - Request

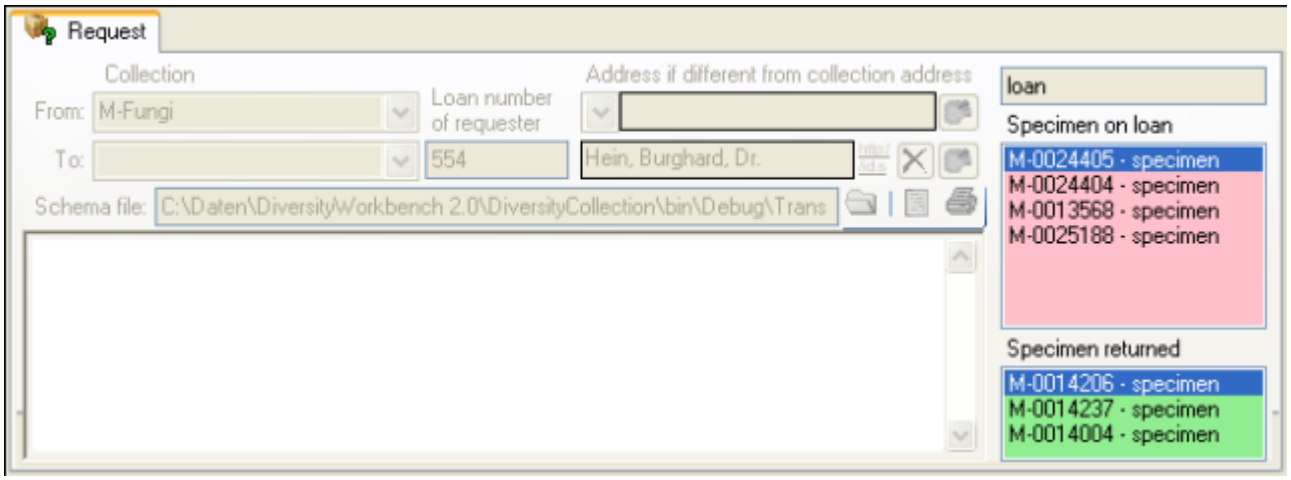
To enter a request for a loan you must be in the user group resp. role DiversityCollectionRequester (see user administration for further details). This is done by the collection managers for their collections. To enable a user to place requests, choose **Administration - Loan requesters** ... from the menu. A window as shown below will open.



As a collection manager you can give requesters access to the collections you manage. Click on the < or > buttons to add or remove collections from the list for which a user can place a request. If the option **Include subcollections** is choosen, a user can request for specimens stored in subcollections of the administrating collection. To give you an overview of the subcollections, the hierarchy of the collection is shown in the tree below the list.

If you are a collection manager and there are requests for your collections, the administration menu will contain a **Loan requests ...** entry. Choose it to open a window listing the requests for specimen in your collections.

If you have entered request for specimen in a foreign collection, the administration menu will contain a **My requests...** entry. To inspect your request choose this entry from the menu to open a window as shown below. The window will show your requests and loans from foreign collections.

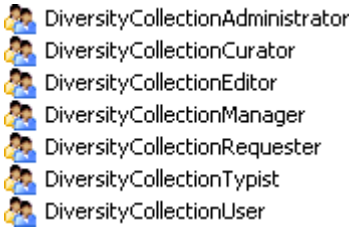


To enter a new request, click on the button. Use the button to search for specimen in the collection and the button to remove unwanted specimen from your list.

Transactions

Prerequisites

The transfer of a specimen between collections is handled with transactions. There are two prerequisites to use transactions in DiversityCollection. You must be in the user group resp. role DiversityCollectionManager (see user administration).

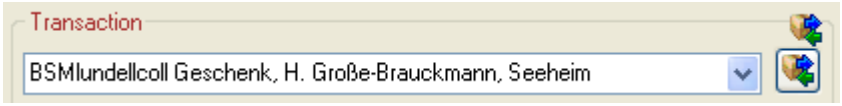


and the collections you have the right to handle transactions for must have been assigned to you as shown in the managers section.

Visibility of transactions

A specimen may be involved in diverse transactions between collections like loan, purchase, gift or exchange. The transactions are shown in the tree for the parts of a specimen and symbolized with an icon. For specimens that are still on loan an icon as shown while a returned loan is symbolized with an icon (see image below).

If you are a collection manager you can take a look at the details of a transaction. Select the specimen part in the tree and then click on the button that will be shown in the details next to the name of the transaction. This will open the window for the transactions as described below.

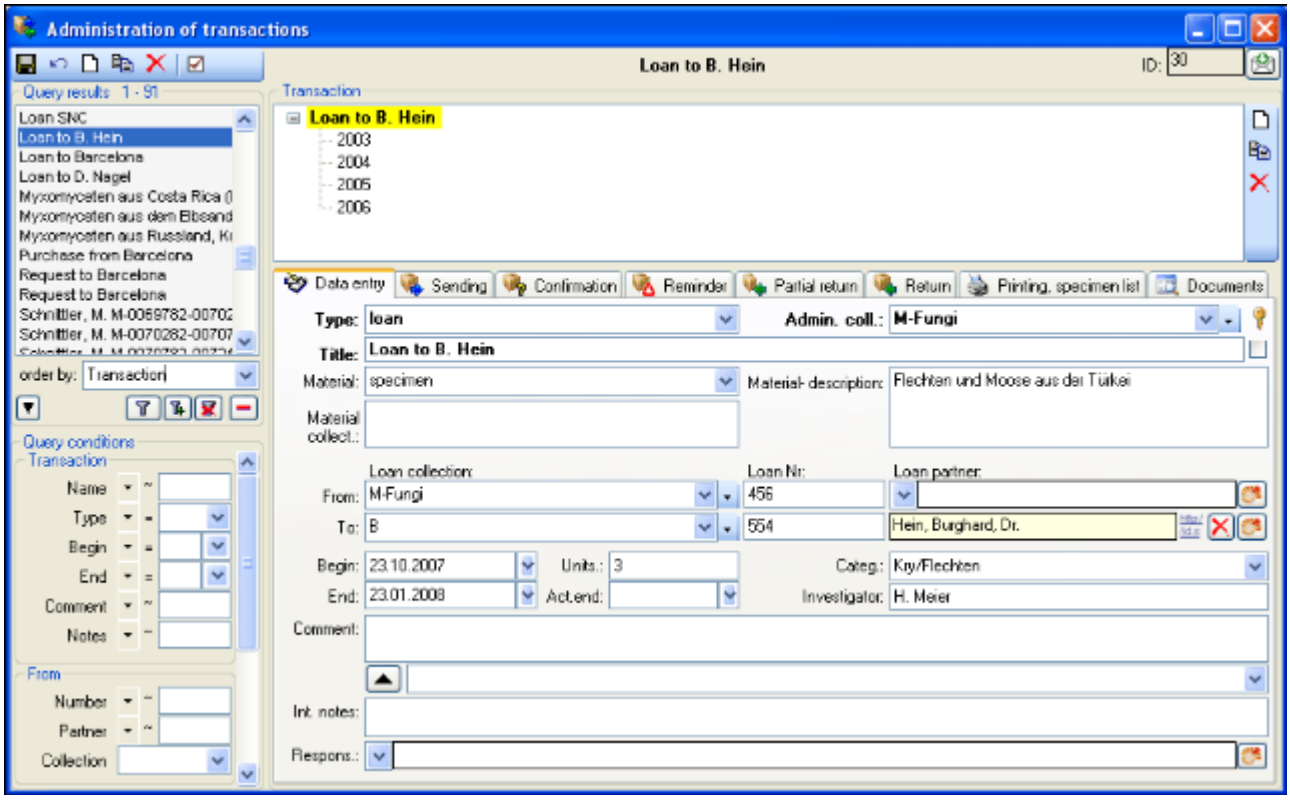


To edit the transactions choose **Administration - Transactions** from the menu. A window will open as shown below. In the tree at the top the relations between the transactions are shown. To create a transaction, dependent on another transaction, choose the superior transaction in the tree and click on the button in the panel right from the tree. With the button you can create a copy of a transaction. To delete a specimen from the list click on the button. The data for the transactions are stored in the tables Transaction and CollectionSpecimenTransaction.

Data entry

On this page you can edit the data of a transaction. Every transaction is linked to an administrating collection (symbolized by the key), and can be edited exclusively by

collection managers of this collection. To appoint managers choose **Administration - Managers** from the menu.

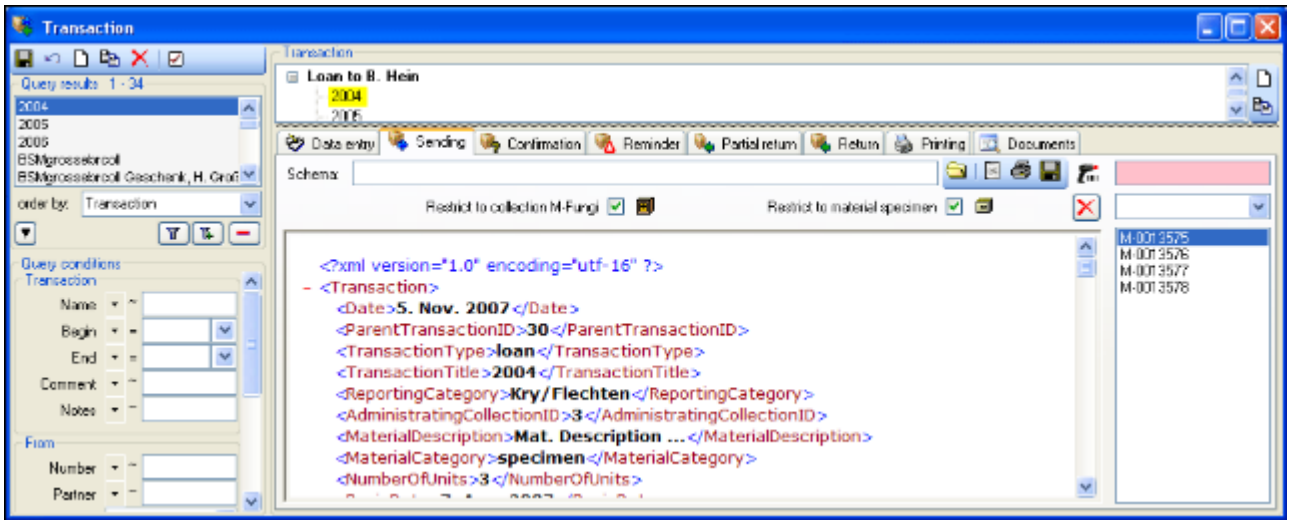


Depending on the type of the transaction additional tab pages will appear for creating documents.

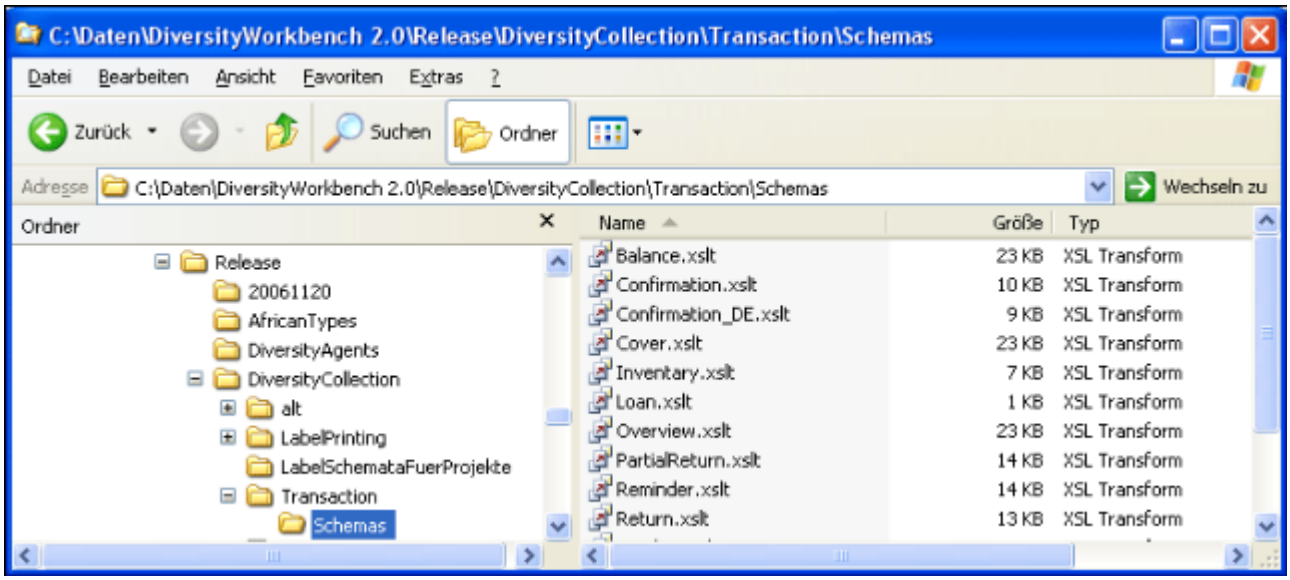
Transaction type	***Tab pages***					
Sending	Loan	Borrow	Exchange	Gift	Purchase	
Confirmation	Loan	Borrow	Exchange	Gift	Purchase	Request
Reminder	Loan	Borrow	Exchange	Gift	Purchase	
Partial return	Loan	Borrow				
Return	Loan	Borrow				
Printing	Loan	Borrow	Exchange	Gift	Purchase	Request
Documents	Loan	Borrow	Exchange	Gift	Purchase	Request
Balance			Exchange			

Common notes

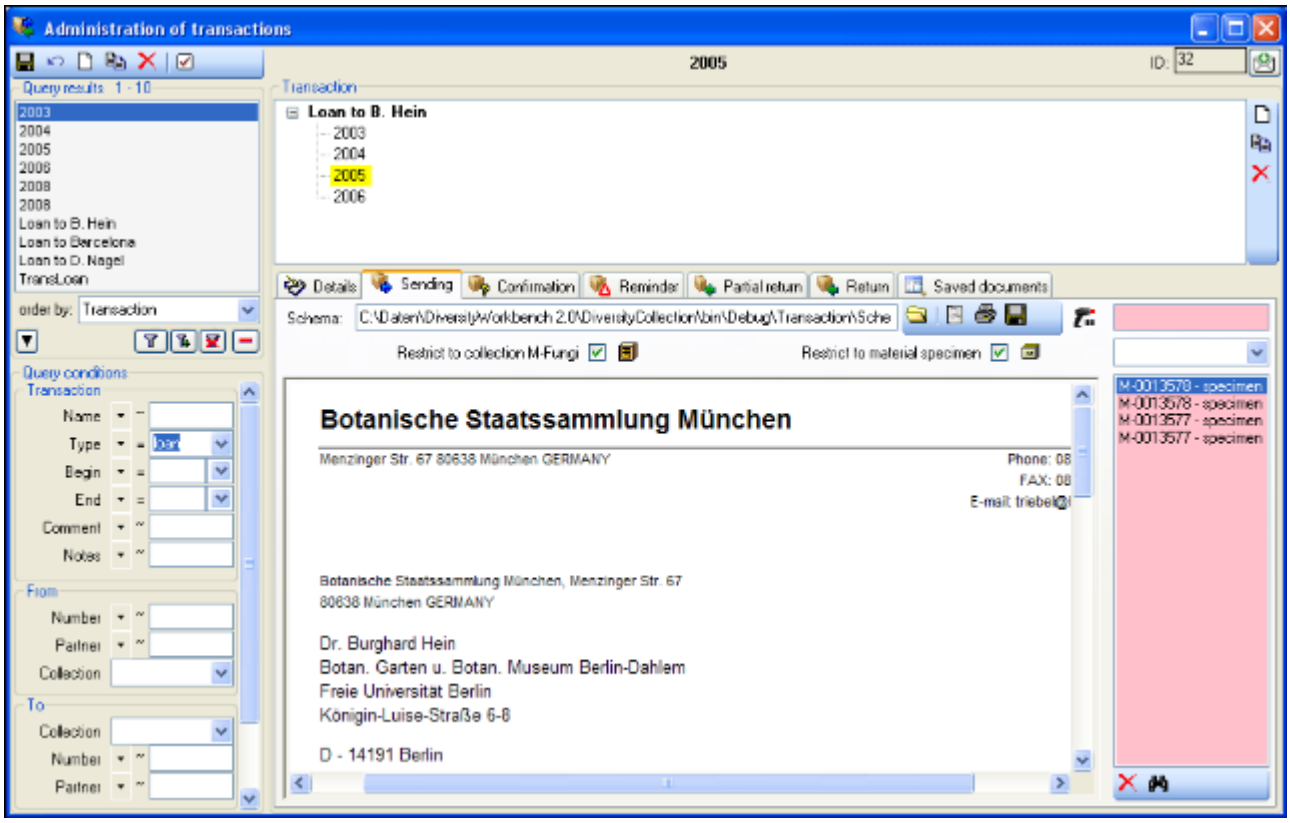
On the tab pages for generating documents you have to specify a xml-schema for your document - click on the button to choose one of the predifined schemas or create your own schema. If you do not specify a schema (i.e. the field for the path of the schema is empty - see below) for your form you will see the XML output created by the database as shown below.



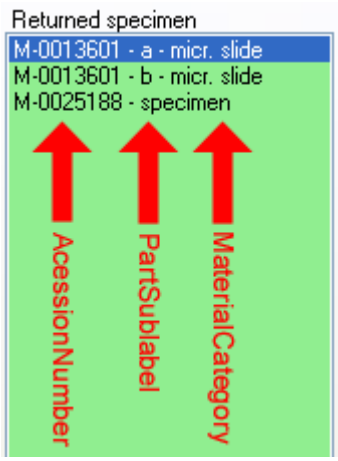
To choose a schema click on the button and select a schema from the list. DiversityCollection contains several ready to use schemas in the folder **Transaction - Schemas**.



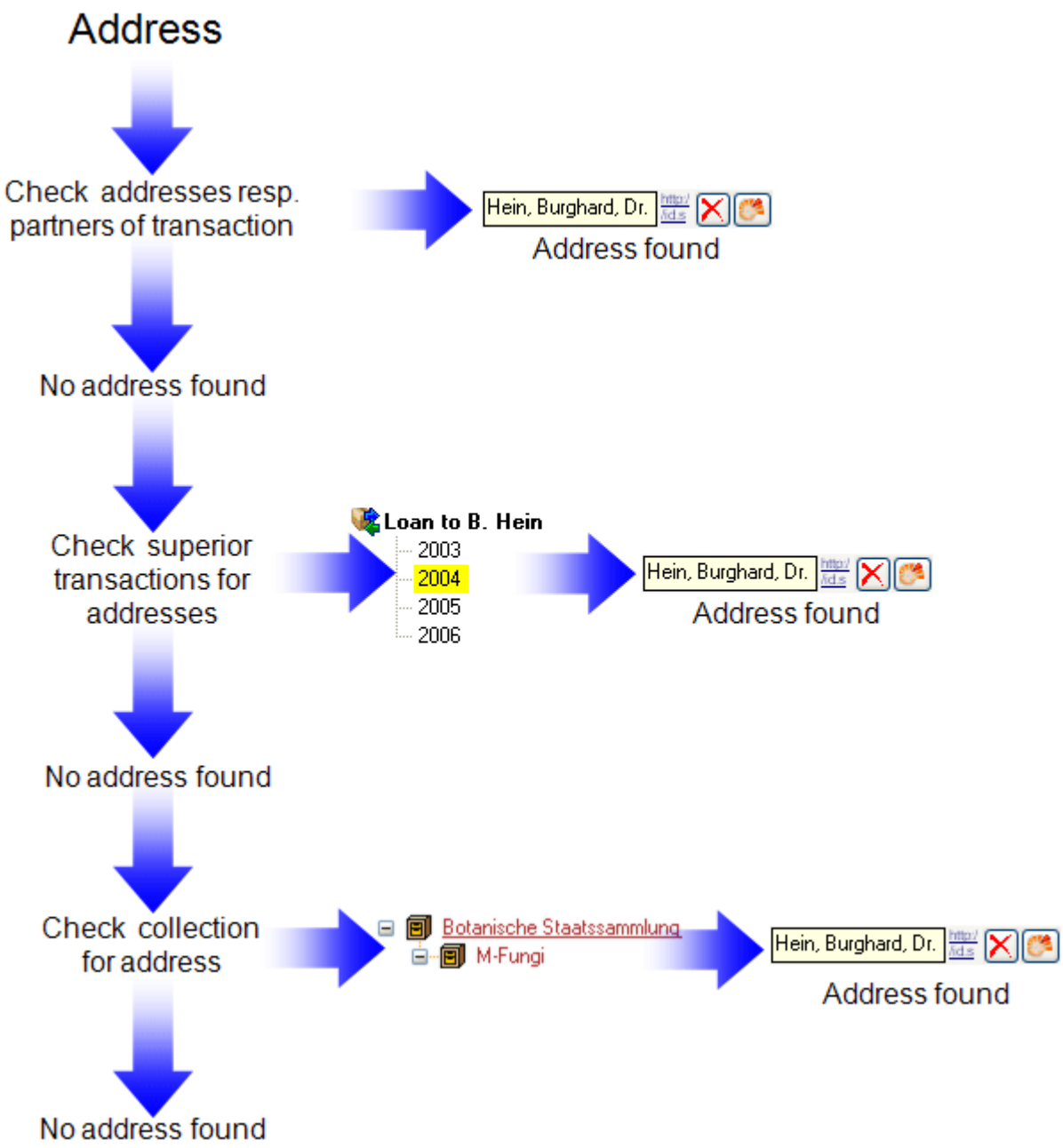
Feel free to change these schemas to your own needs (editors are available e.g. from http://www.altova.com/). Choose the schema you need and then click on the button to create a document. To print the document, use the button. To store the current document in the documents for later reference click on the button. An example for a document is shown below.



The specimens will be listed with their accession number, a optional label of the part and the material category to ensure discrimination in case a specimen contains more then one part (see image below).



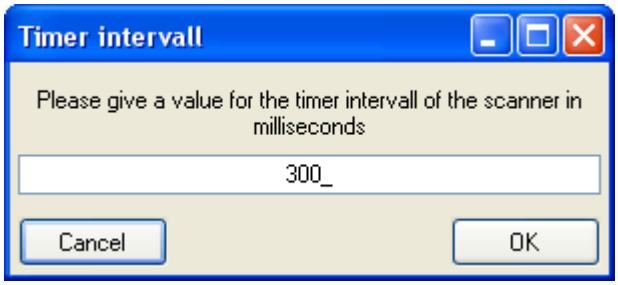
If a transaction entry has no address, resp. a transaction partner linked to an entry in the module DiversityAgents, where addresses are stored, the program will try to find an address in one of the superior transactions from the transaction partners found there. If no address exists in these transactions, the program will try to get an address from the collections. Therefore a transaction partner linked to DiversityAgents is only necessary if either there are no addresses for the collections or you want to use a different address. The search path as described is summarized in the image below.


Address
Check addresses resp. partners of transaction
Hein, Burghard, Dr.
Address found
No address found
Check superior transactions for addresses
Loan to B. Hein
2003
2004
2005
2006
Hein, Burghard, Dr.
Address found
No address found
Check collection for address
Botanische Staatssammlung
M-Fungi
Hein, Burghard, Dr.
Address found
No address found

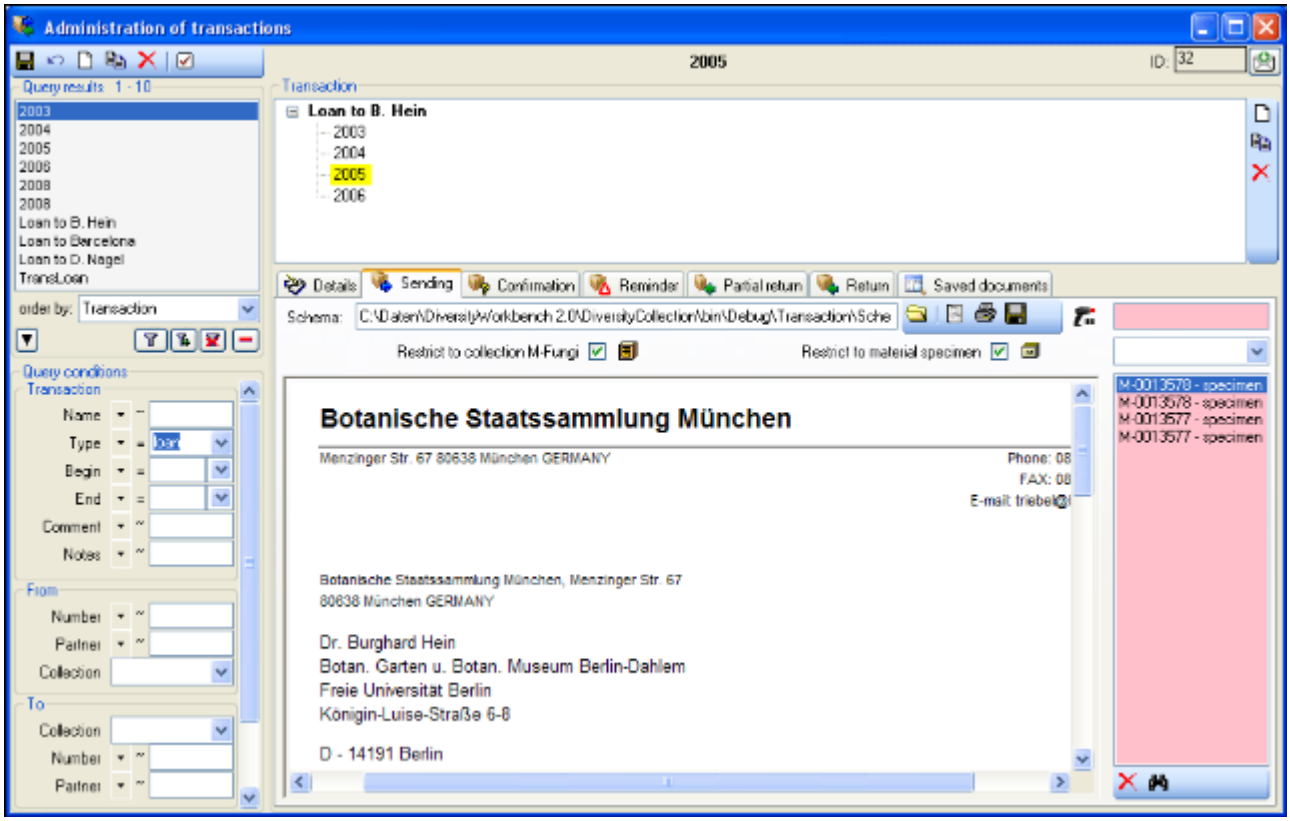
Transactions - Sending

In the sending tab page of the transaction window you can create the covering note for a sent sample. To choose a schema click on the button and select a schema from the list. DiversityCollection contains several ready to use schemas in the folder **Transaction - Schemas**.

To add a specimen to the list, you can either use the combobox or a barcode scanner. In case you scanner is reading only parts of the barcode try to adjust the timer intervall - click on the button to open a window as shown below.



To restrict the selection of the specimens and parts of the specimens, you can check the **Restrict to collection** ... and **Restrict to material** ... options. To remove a specimen from the list, use the button under the specimen list. If you want to see the details of a specimen, choose it in the list and click on the button. Click on the button to create the document as shown below.



Transactions - Confirmation

On this page you can create an inquiry letter for a package sent, e.g. if a parcel was sent to a loan taker and no confirmation that the parcel reached its destination was returned so far.

With the button, choose the schema you need. Click on the button to create a document. To print the document, use the button. To store the current document in the documents for later reference click on the button.

Transactions - Reminder

If a loan is due to return resp. a loan taker did not meet the deadline, you can create a prompt note here.

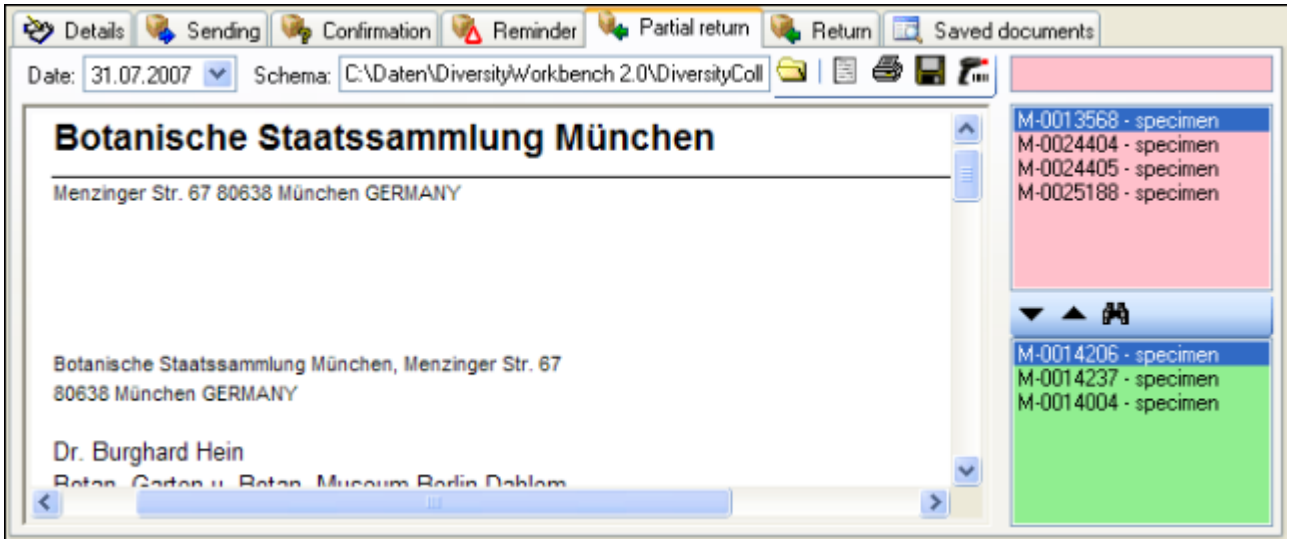
With the button, choose the schema you need. Click on the button to create a document. To print the document, use the button. To store the current document in the documents for later reference click on the button.

The at the upper right the missing specimen and at the lower right the returned specimen are listed.

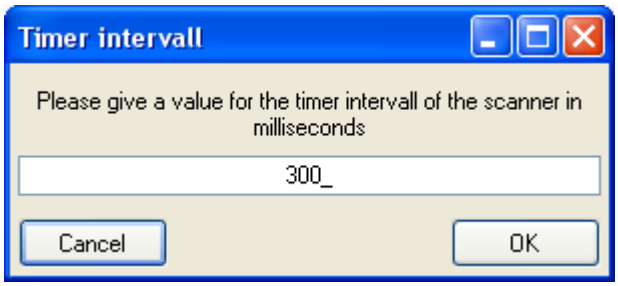
Transactions - Partial Return

If parts of a loan were returned, create a letter of acknowledgment here.

With the button, choose the schema you need. Click on the button to create a document. To print the document, use the button. To store the current document in the documents for later reference click on the button.

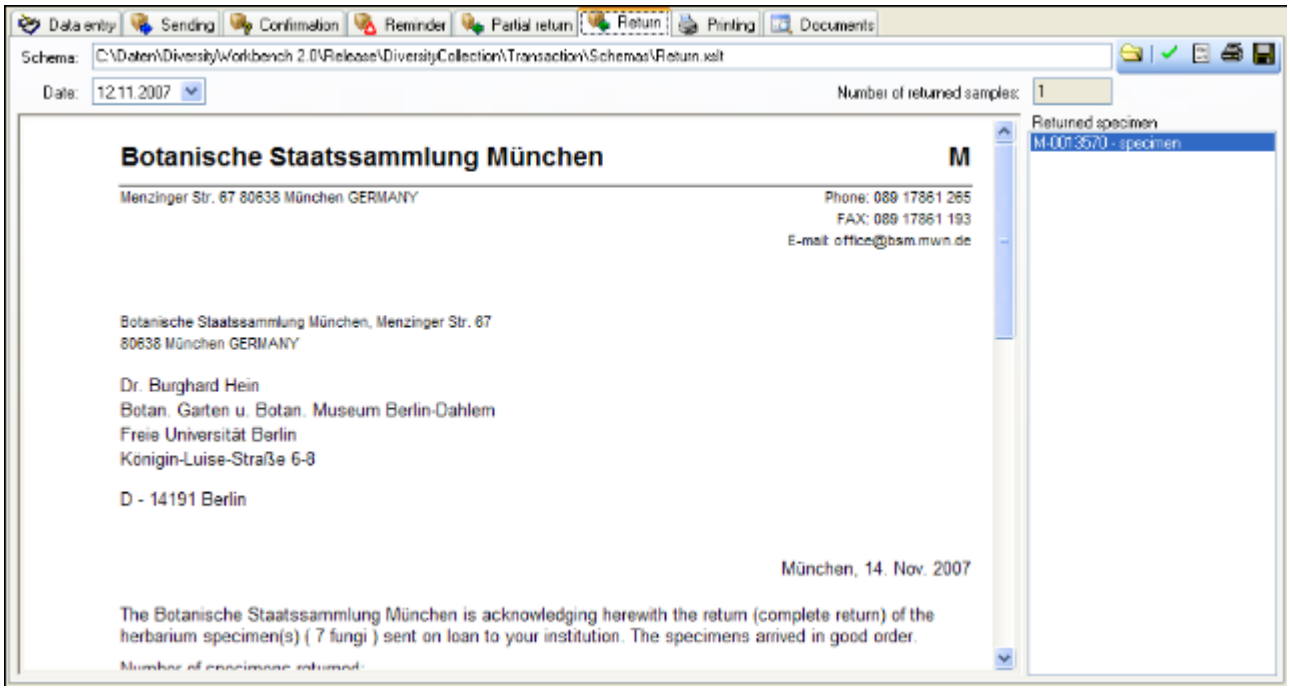


You have two option to enter returned specimen. Use the ▲and ▼buttons respectively to move items between the lists or use a barcode scanner - move the mouse cursor into the field for the barcode detection, then scan the barcode. The specimen will be automatically inserted into the list for the returned specimen. In case your scanner is reading only parts of the barcode try to ajust the timer intervall - click on the button to open a window as shown below. Here you can set the interval to a value that is compatible to your scanner. If you want to see the details of a specimen, choose it in the list and click on the button.



Transactions - Return

If a loan is returned, create a letter of acknowledgment here. With the button, choose the schema you need. If all specimen were returned, click on the button. This take all remaining specimen into the list of the returned specimens. With the date field you can define a different date as the start if e.g. you starting to register the returned specimen some days before. Click on the button to create a document. To print the document, use the button. To store the current document in the documents for later reference click on the button.

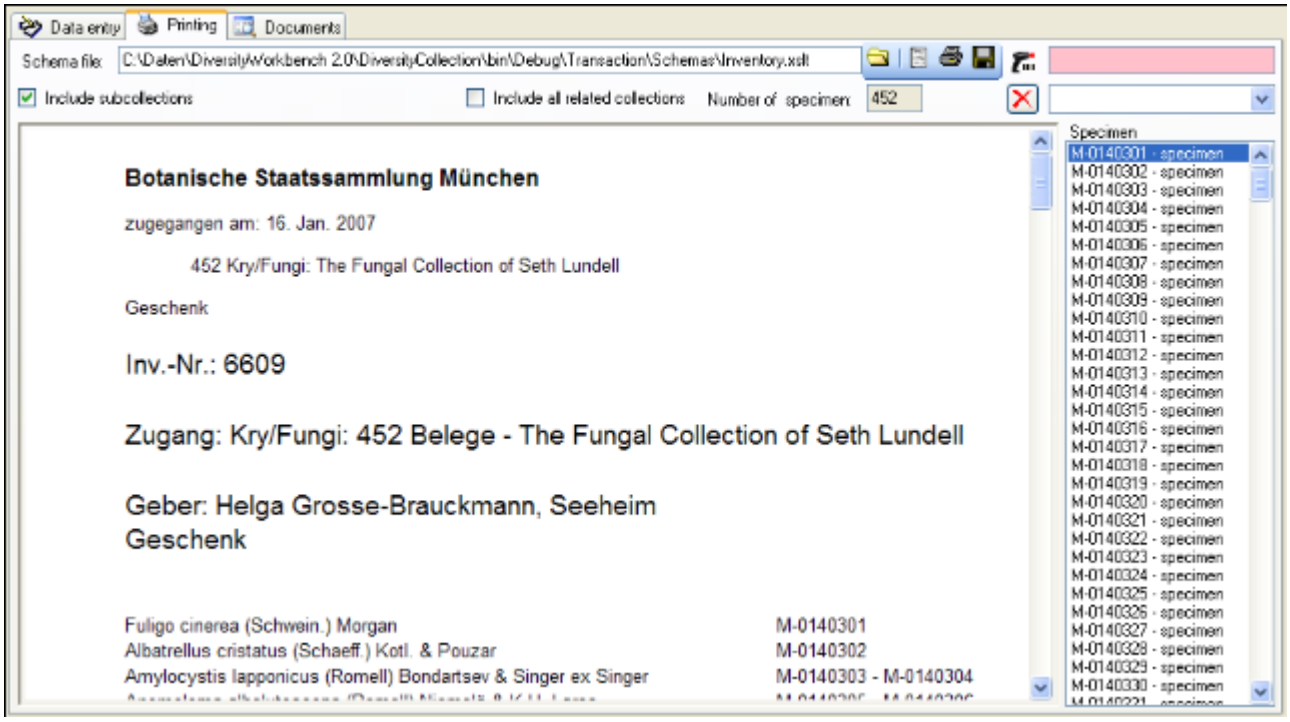


Transactions - Printing / Inventory

Besides the special pages for creating letters along with a loan this page provides the possibility to print cover letters, inventories and the like. You can restrict the selected specimen to the current collection, include the subcollections or all related collections - just activate the corresponding checkboxes.

To add a specimen to the list, you can either use the combobox or a barcode scanner. To use the barcode scanner, move your mouse cursor into the pink field right from the scanner. In case your scanner is reading only parts of the barcode try to adjust the timer intervall - click on the button to open a window where you can set the timer intervall to a different value. To remove specimen from the list, select it and then click on the button.

With the button, choose the schema you need. Click on the button to create a preview of the document. To print the document, use the button. To store the current document in the documents for later reference click on the button.



Transactions - Documents

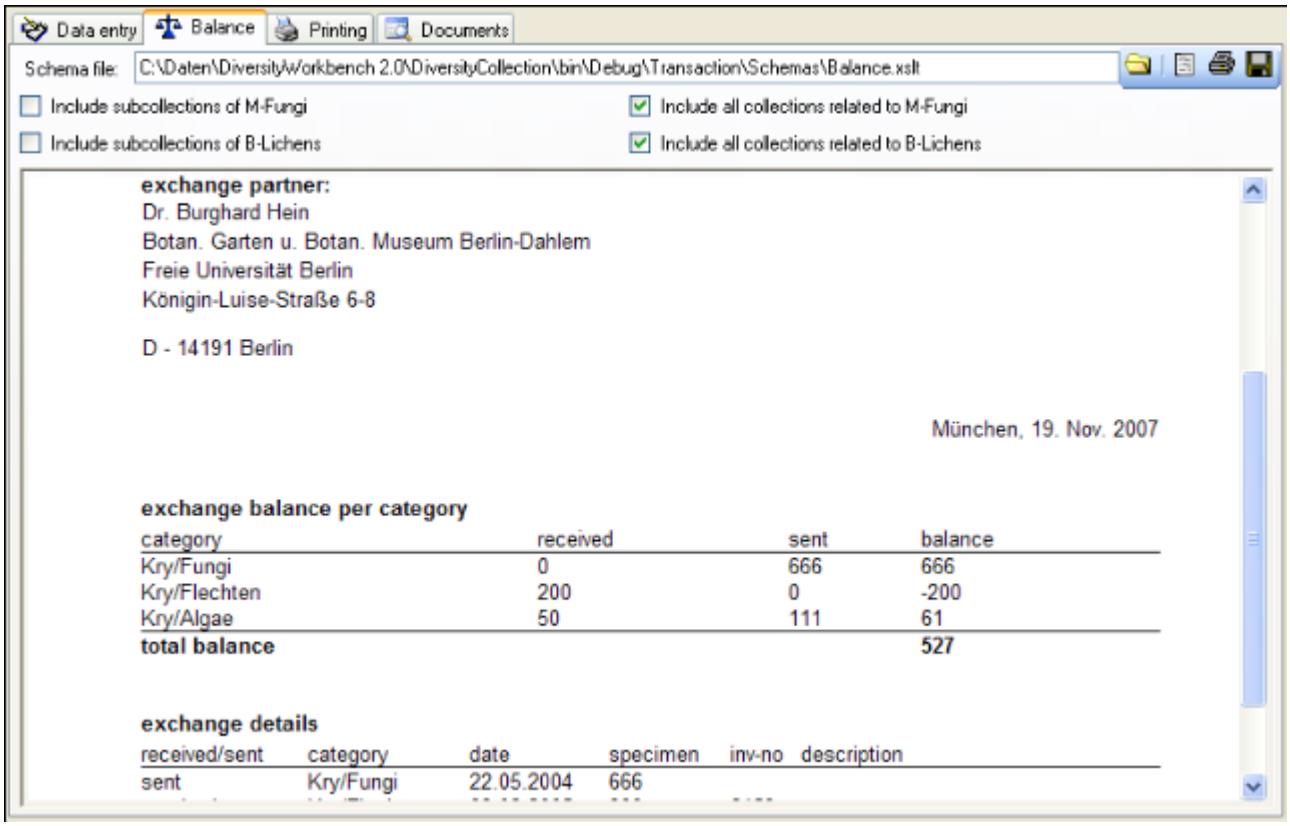
This page stores all the documents created or received along with a transaction.

To add a document, scan the document and create a screeshot of this document, then create a new entry (click on the button) and insert the screenshot with the Add image from document button. In the lower part you can enter any text related to the document.

Transactions - Balance

The balance for the exchange between two collections.

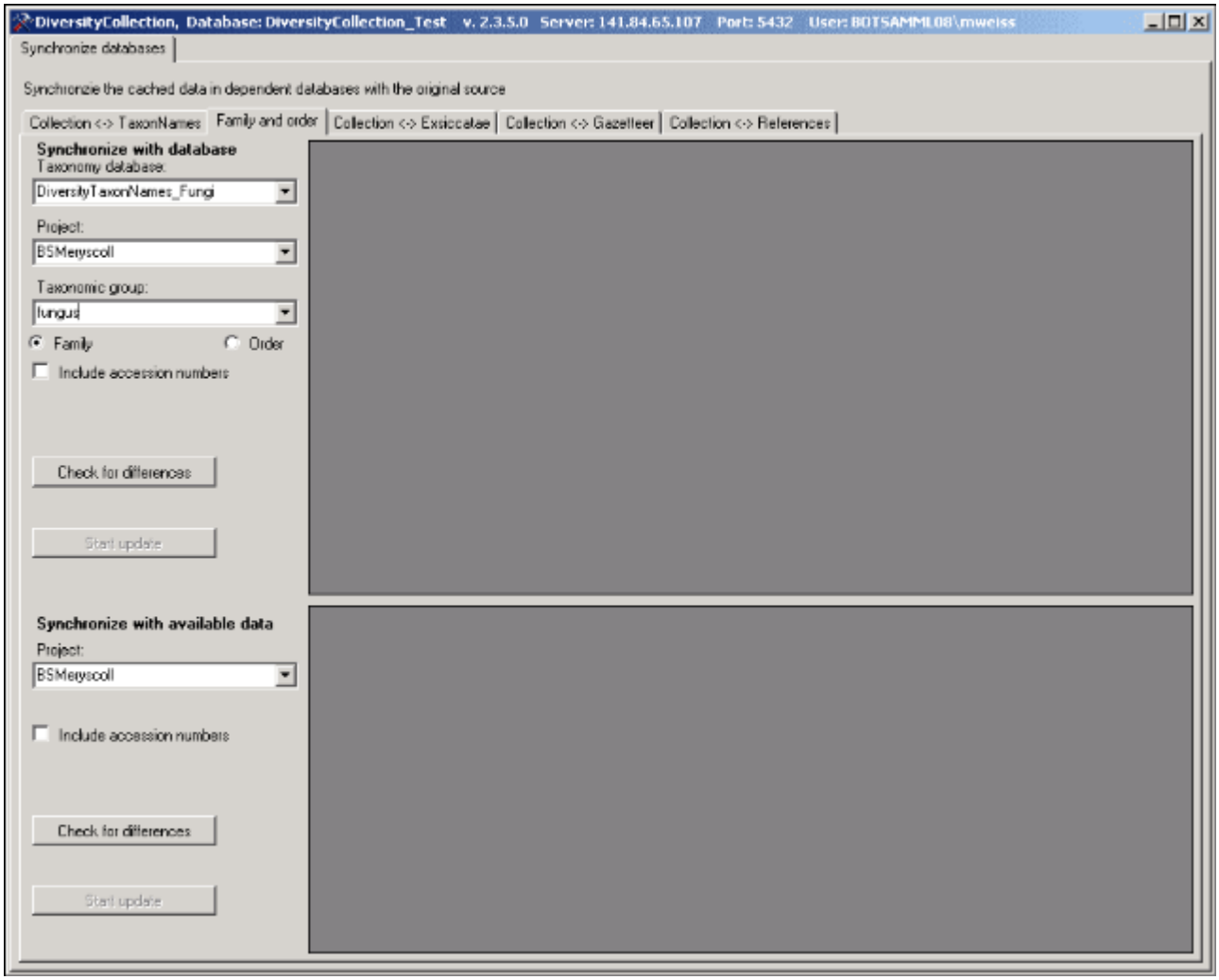
With the button, choose the schema you need. Click on the button to create a document. To print the document, use the button. To store the current document in the documents for later reference click on the button. To include either the subcollections of any related collections to the collections of the current transaction select the corresponding checkboxes.



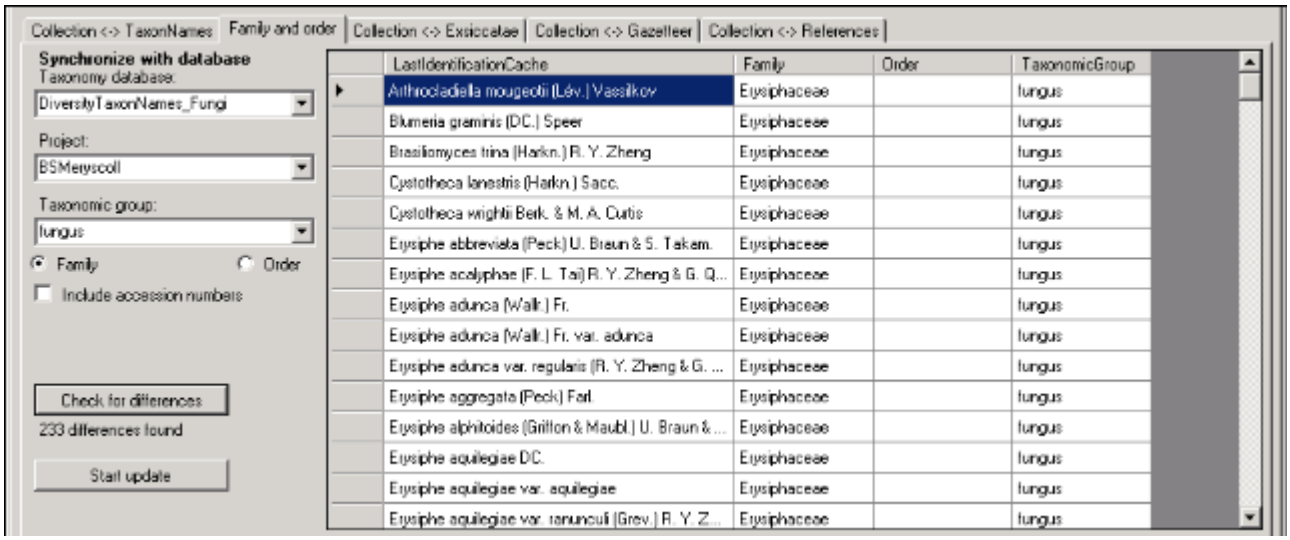
Maintenance

DiversityCollection can be linked with several other modules of the Diversity Workbench. If you link a data source to another module, the URI of the dataset in the other module together with several cached values will be stored in DiversityCollection. Along with changes in the other modules these cached values may differ from the original values. To get the actual values you can use the maintenance function within DiversityCollection.

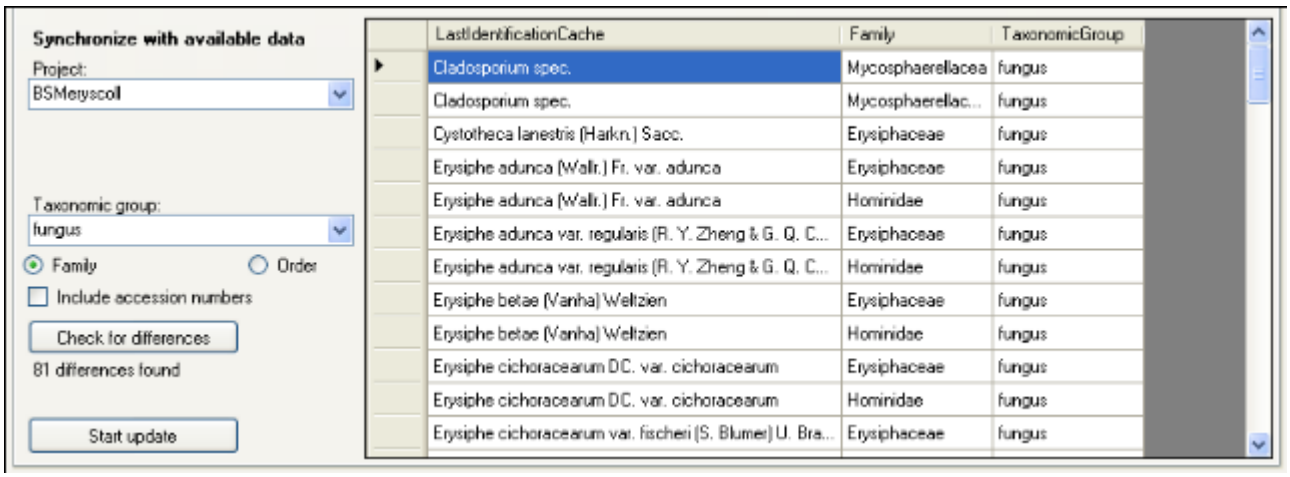
Choose **Administration ->** **Maintenance ...** from the menu. A window as shown below will open. To transfer the higher taxonomic entries choose the tab Family and order. The upper part will synchronize the entries with taxonomic databases like DiversityTaxonNames_Fungi. Select a taxonomic database, a project and a taxonomic group, then choose if you want to synchronize the family or the order. If you want to take a look at the single datasets you have to select the "Include accession number" checkbox. Than start the query with a click on the Check for differences button.



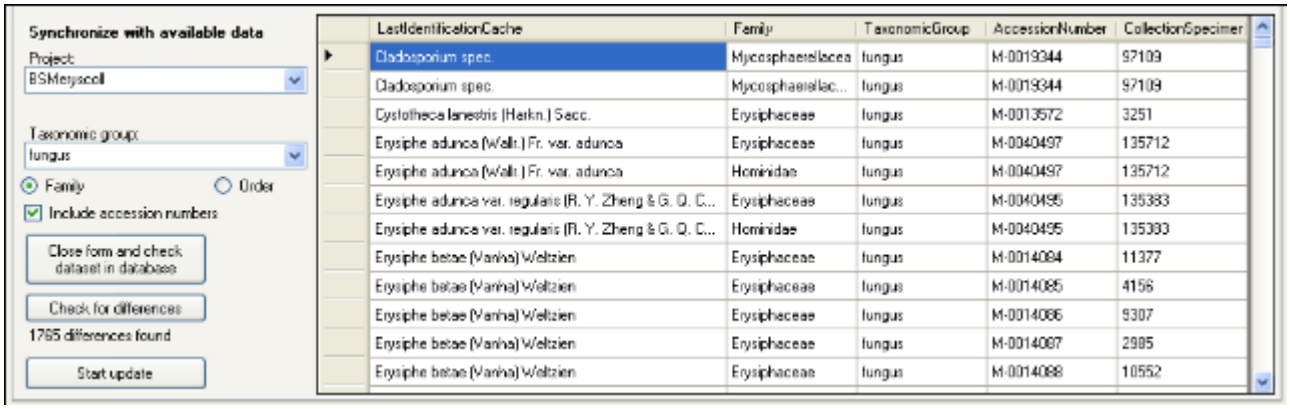
The form will list all differences found as shown below. To update the database click on the Start update button.



In the lower part you can synchronize your entries within the database. Choose a project and click on the Check for differences button to start the query. To import the higher taxa to the dataset click on the Start update button.



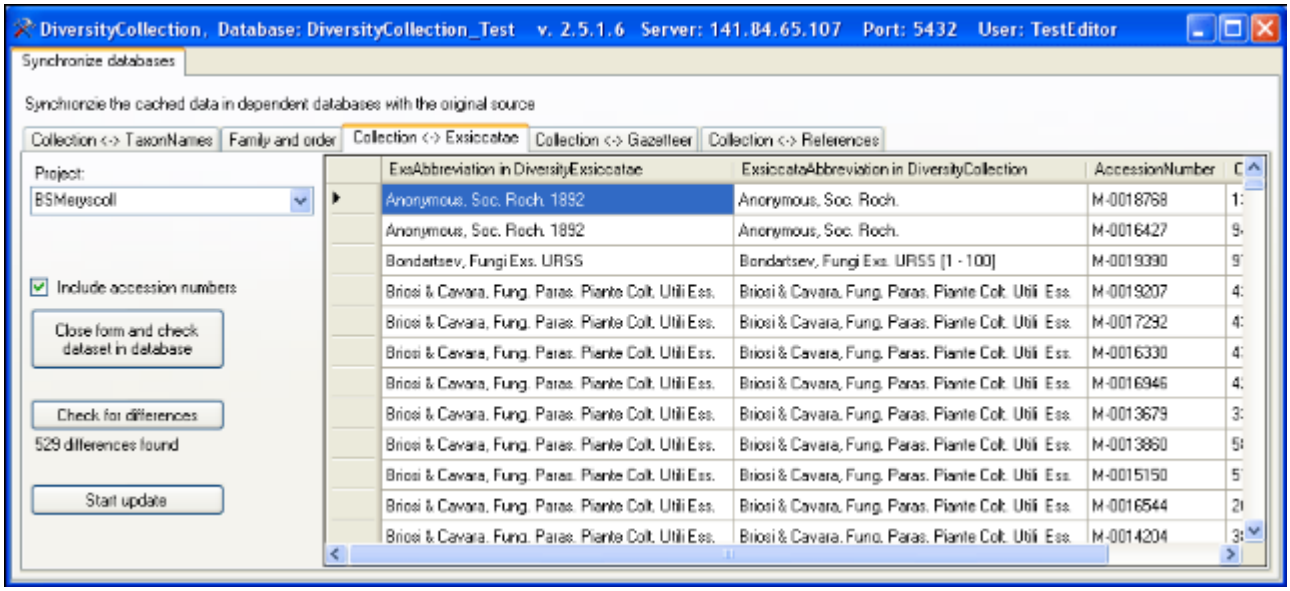
If you want to have a more detailed look on your data you have to check the "Include accession number" checkbox before starting the query. A button Close form and check dataset in database will appear that will take you back to a single dataset in the database.



Maintenance - Exsiccatae

To synchronize the abbreviations of the titles of the exsiccatal series that are linked to the module DiversityExsiccatae choose **Administration -> Maintenance...** from the menu. A window as shown below will open. On the tab page **Collection <-> Exsiccatae** select the project for which the titles should be synchronized. The title is stored in the field ExsiccataAbbreviation in the table CollectionSpecimen. Select the project that should be synchronized. Then start the query with a click on the

button.



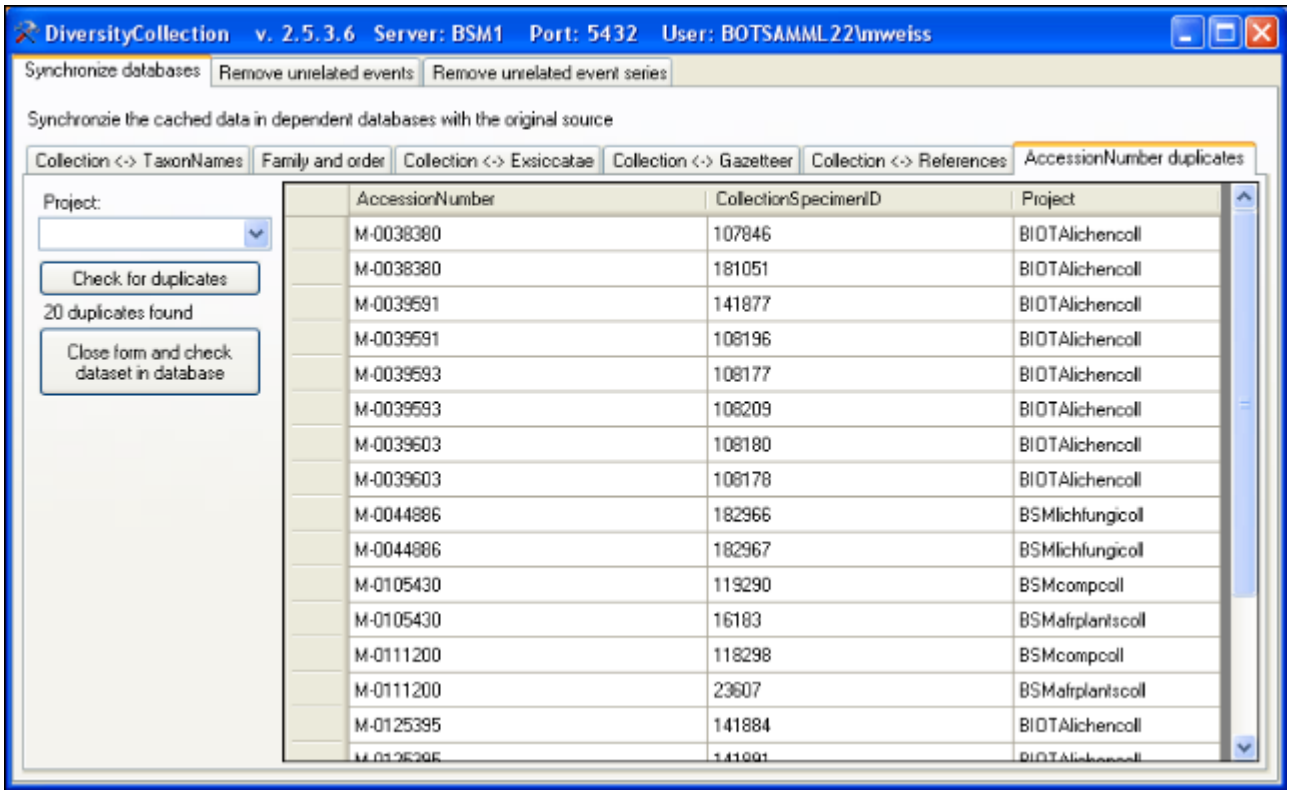
The form will list all differences found. To update the database click on the

button. If you want to have a more detailed look at your data you have to check the "Include accession number" checkbox before starting the query. A button

will appear that will take you back to a single dataset in the database.

Maintenance - Accession number duplicates

The database will not prevent you from entering an accession number several times e.g. via an import. To check for duplicate accession numbers choose **Administration -> Maintenance** from the menu. A window as shown below will open. On the tab page **AccessionNumber duplicates** . You can restrict the query to a project. To start the query with a click on the **Check for duplictes** button.



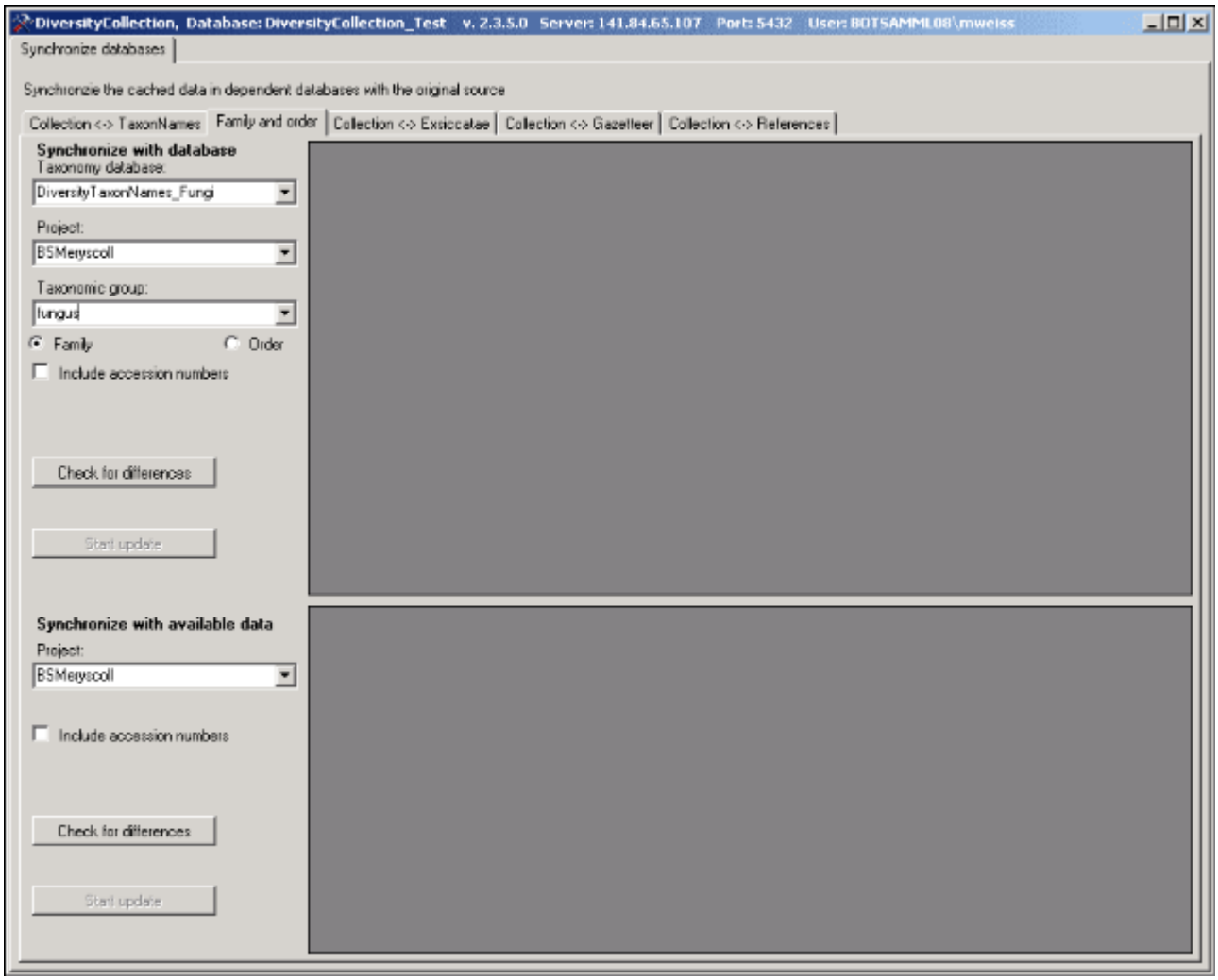
The form will list all duplicates found as shown above. To switch to a dataset in the database, select it in the table and click on the

button.

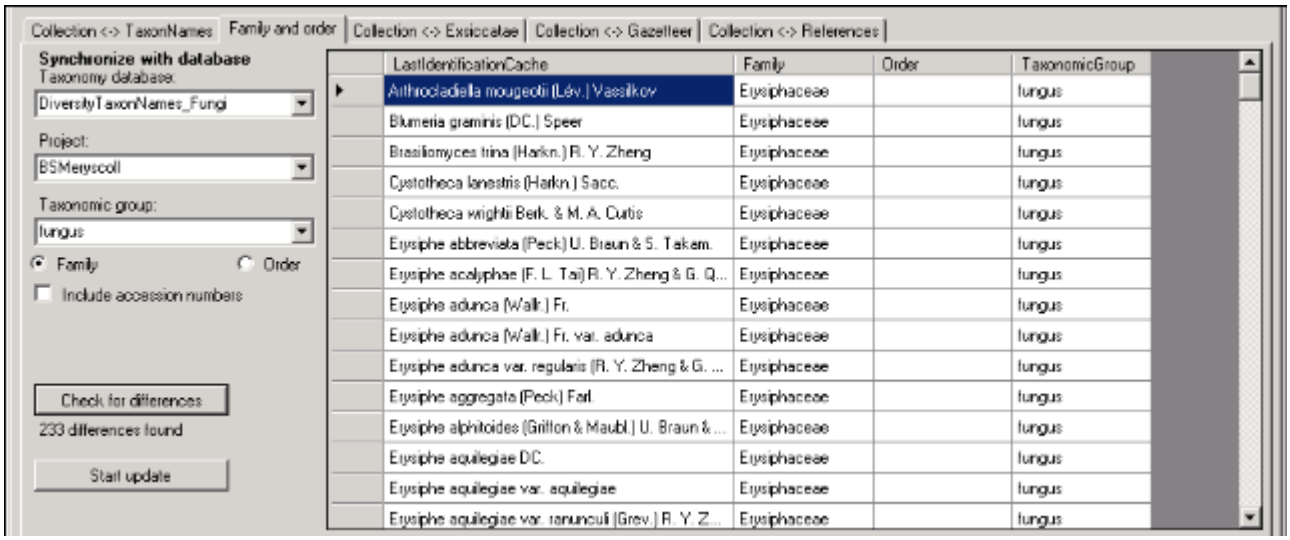
Maintenance - Family and Order

To synchronize the entries for the family and order of taxa derived from the module DiversityTaxonNames choose **Administration -> Maintenance** from the menu. A window as shown below will open. To synchronize the higher taxonomic entries for entries linked to the module DiversityTaxonNames choose the tab **Family and order**. These are stored in the fields FamilyCache and OrderCache in the table IdentificationUnit .

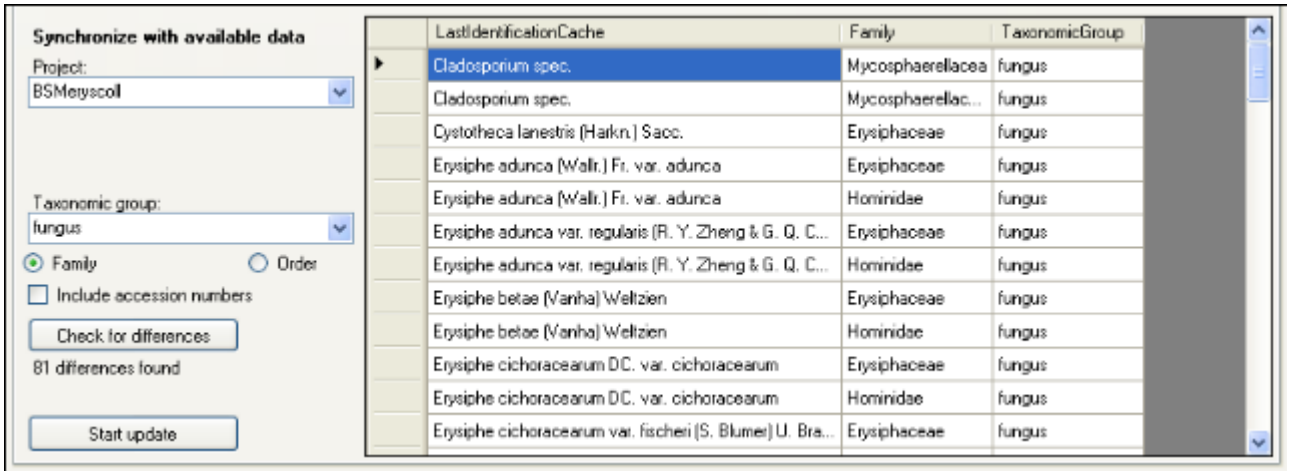
In the upper part you can synchronize your data with the entries in taxonomic databases like DiversityTaxonNames_Fungi. Select a taxonomic database, a project and a taxonomic group, then choose whether you want to synchronize the family or the order. If you want to inspect single datasets you have to check the **Include accession number** checkbox. Then click on the Check for differences button to start the query.



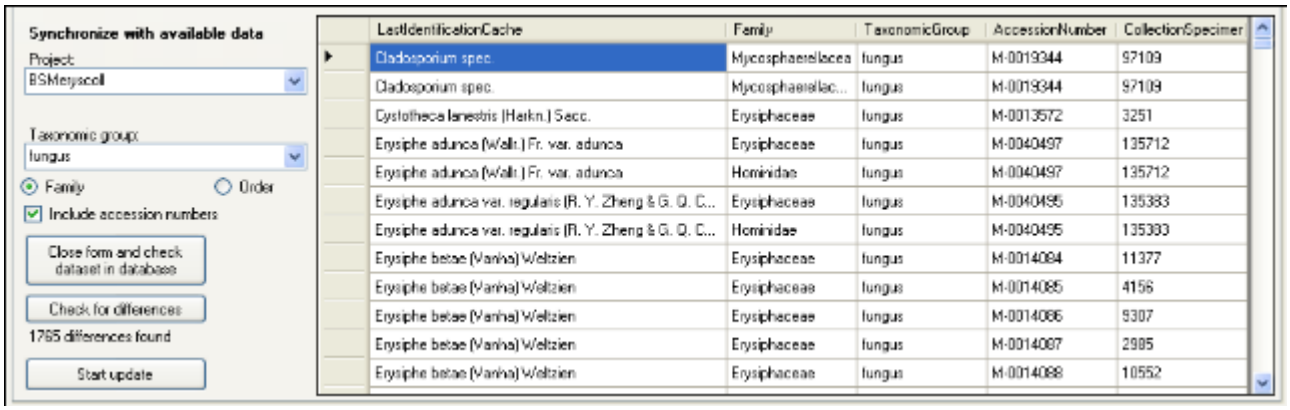
The form will list all differences found as shown below. To update the database click on the Start update button.



In the lower part you can synchronize your entries within the database. Choose a project, the taxonomic group and if you want to check the family or the order and click on the Check for differences button to start the query. To import the higher taxa to the dataset click on the Start update button.

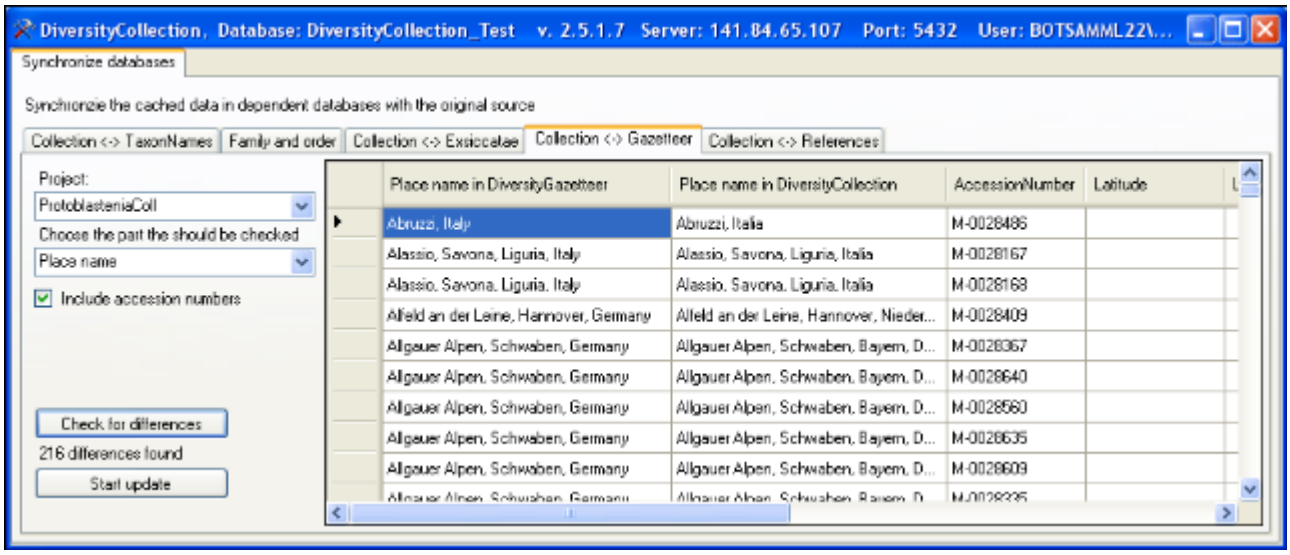


If you want to have a more detailed look on your data you have to check the **Include accession number** checkbox before starting the query. A button Close form and check dataset in database will appear that will take you back to a single dataset in the database.


Synchronize with available data
Project:
BSMeryscoll
Taxonomic group:
fungus
Family
Order
Include accession numbers
Close form and check dataset in database
Check for differences
1765 differences found
Start update
LastIdentificationCache
Family
TaxonomicGroup
AccessionNumber
CollectionSpecimer
Cladosporium spec.
Mycosphaerellacea
fungus
M-0019344
97109
Cladosporium spec.
Mycosphaerellac...
fungus
M-0019344
97109
Cystotheca lanestris (Harkn.) Sacc.
Erysiphaceae
fungus
M-0013572
3251
Erysiphe adunca (Wallr.) Fr. var. adunca
Erysiphaceae
fungus
M-0040497
135712
Erysiphe adunca (Wallr.) Fr. var. adunca
Hominidae
fungus
M-0040497
135712
Erysiphe adunca var. regularis (R. Y. Zheng & G. Q. C...
Erysiphaceae
fungus
M-0040495
135383
Erysiphe adunca var. regularis (R. Y. Zheng & G. Q. C...
Hominidae
fungus
M-0040495
135383
Erysiphe betae (Vanha) Weltzien
Erysiphaceae
fungus
M-0014084
11377
Erysiphe betae (Vanha) Weltzien
Erysiphaceae
fungus
M-0014085
4156
Erysiphe betae (Vanha) Weltzien
Erysiphaceae
fungus
M-0014086
9307
Erysiphe betae (Vanha) Weltzien
Erysiphaceae
fungus
M-0014087
2985
Erysiphe betae (Vanha) Weltzien
Erysiphaceae
fungus
M-0014088
10552

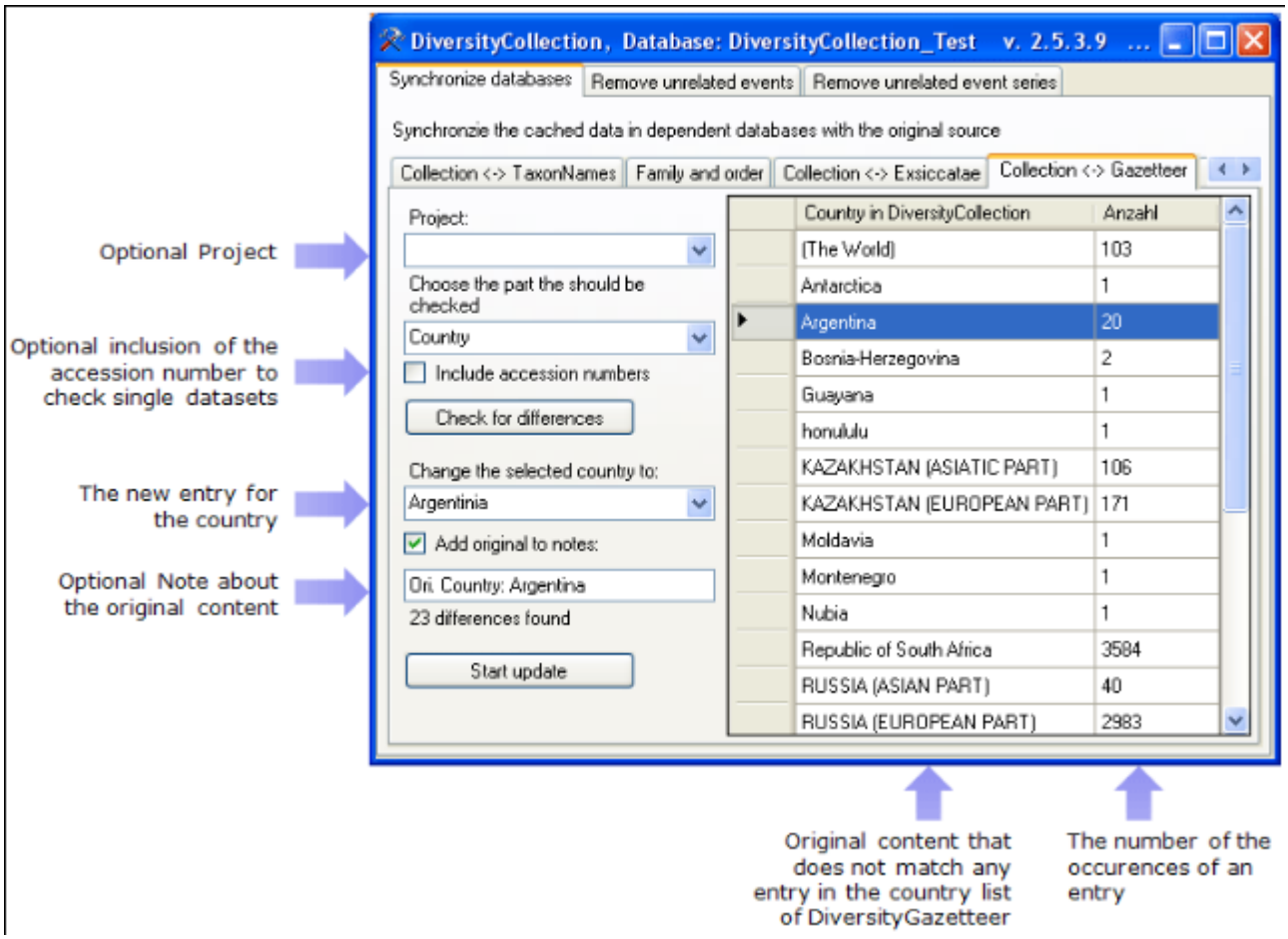
Maintenance - Gazetteer

To synchronize the entries derived from the module DiversityGazetteer choose **Administration -> Maintenance** from the menu. A window as shown below will open. On the tab page **Collection <-> Gazetteer** select the project for which the entries should be synchronized. There are 3 targets for the synchronization: Place name in the field Location1 in table CollectionEventLocalisation, Country stored in the field CountryCache in the table CollectionEvent and the Coordinates stored in the fields AverageLatitudeCache and AverageLongitudeCache in the table CollectionEventLocalisation. Select one of there targets for the synchronization. To start the query, click on the Check for differences button.



The form will list all differences found. To update the database click on the Start update button. If you want to have a more detailed look on your data, you have to check the **Include accession number** checkbox before starting the query. A button Close form and check dataset in database will appear that will take you back to a single dataset in the database.

To check for countries that do not match entries in DiversityGazetteer choose Country from the list as shown below.

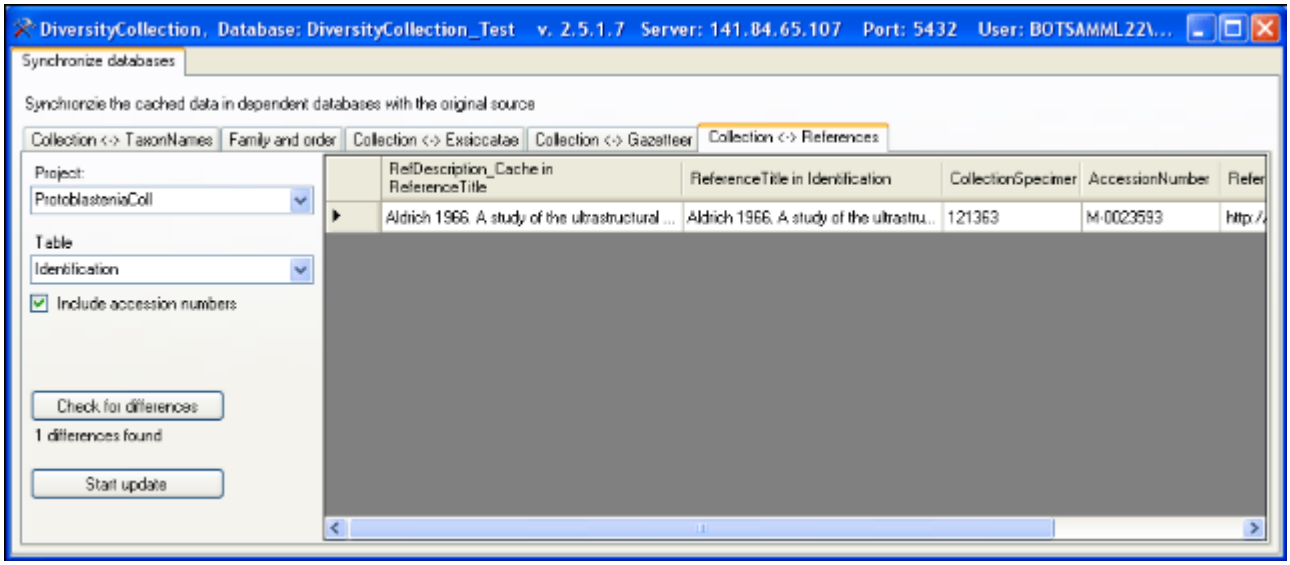


The image above summarizes the options for an update of the country. You can restrict your query to one of the projects you have access to. If you want to check single datasets, check the **include accession numbers** option before you start the query. To keep the old entry in the Notes field check the **Add original to notes** option.

Maintenance - References

To synchronize the titles of the references that are linked to the module DiversityReferences choose **Administration -> Maintenance** from the menu. A window as shown below will open. On the tab page **Collection <-> References** select the project for which the reference titles should be synchronized. There are 3 tables which may contain links to DiversityReferences: CollectionEvent, CollectionSpecimen and Identification. Choose one of these tables for the synchronization. Then start the query with a click on the

button.



The form will list all differences found. To update the database click on the

button. If you want to have a more detailed look on one of the datasets in the list data you have to check the "Include accession number" checkbox before starting the query. A button

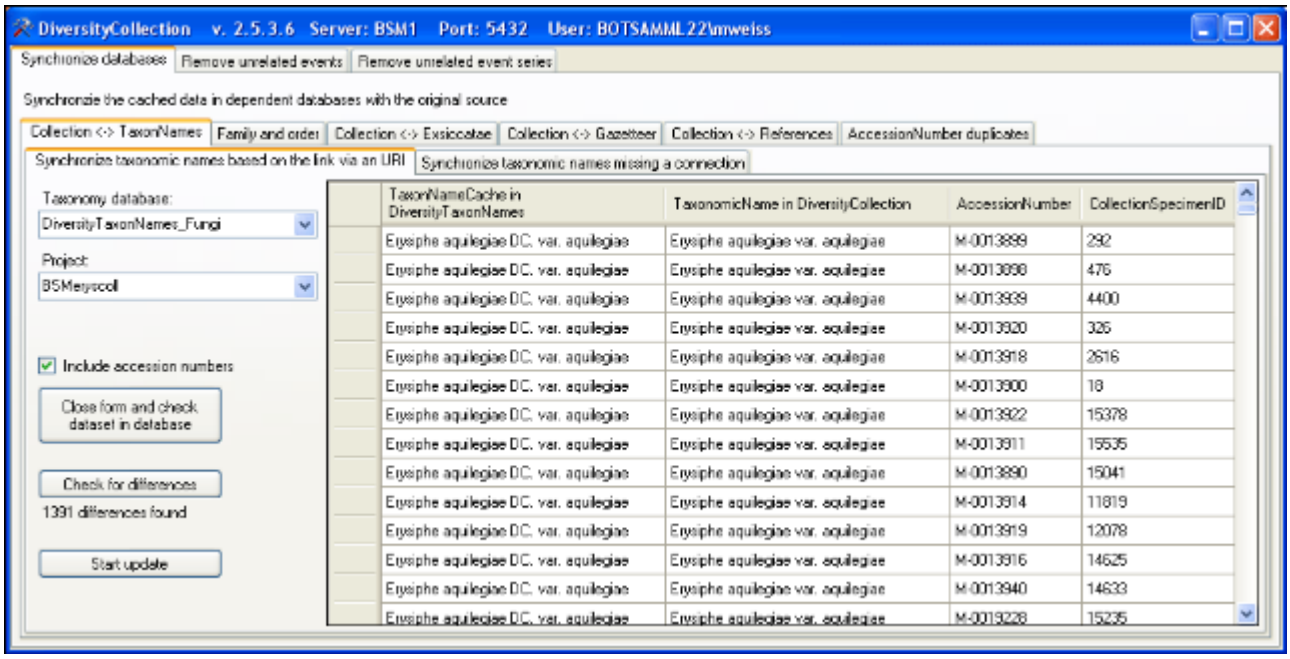
will appear that will take you back to a single dataset in the database.

Maintenance - Taxonomic names

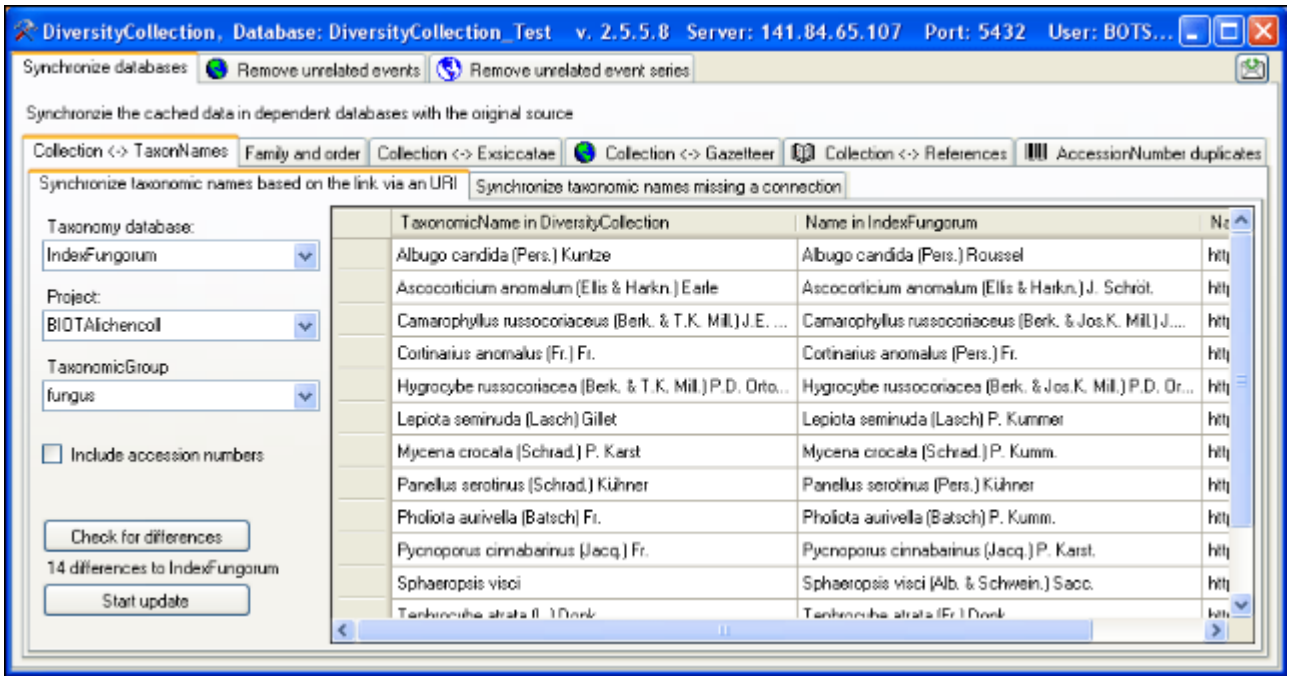
To synchronize the entries for the taxonomic names derived from the module DiversityTaxonNames choose **Administration -> Maintenance...** from the menu. A window as shown below will open. To synchronize the taxonomic names for entries linked to the module DiversityTaxonNames choose the tab **Collection <-> TaxonNames**. There are two ways to synchronize taxonomic names. You can either **Synchronize taxonomic names based on the link via an URI** for entries where the link to a taxonomic database is allready established or you can **Synchronize taxonomic names missing a connection**, where no link is established and you can query for identical names in one of the databases.

Synchronize taxonomic names based on the link via an URI

The taxonomic names are stored in the field TaxonomicName in the table Identification. Select a taxonomic database and a project that should be synchronized. Than start the query with a click on the Check for differences button.



If you synchronize your data with a webservice as shown in the image below, you can specify the taxonomic group as well.



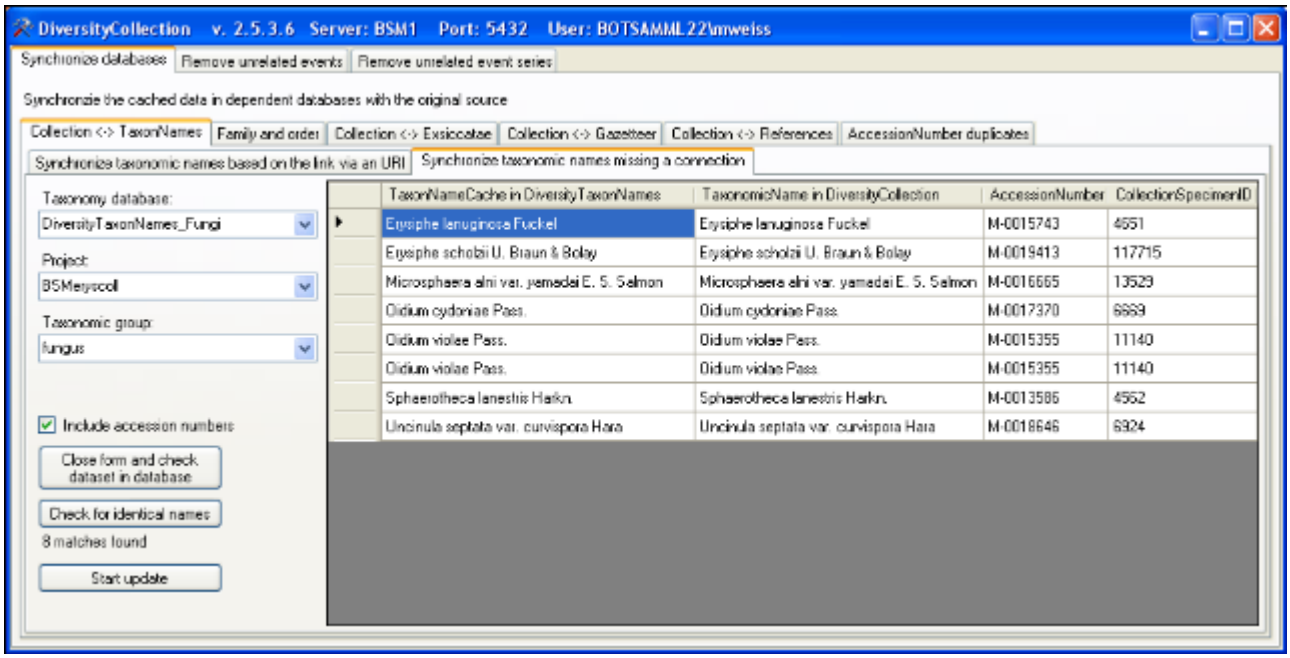
The form will list all differences found. To update the database click on the

button. If you want to have a more detailed look on your data you have to check the **Include accession number** checkbox before starting the query. A button

will appear that will take you back to a single dataset in the database.

Synchronize taxonomic names missing a connection

Select a taxonomic database, a project and a taxonomic group to search for identical names. Than start the query with a click on the **Check for identical names** button.



The form will list all matches found. To insert the links to the database click on the

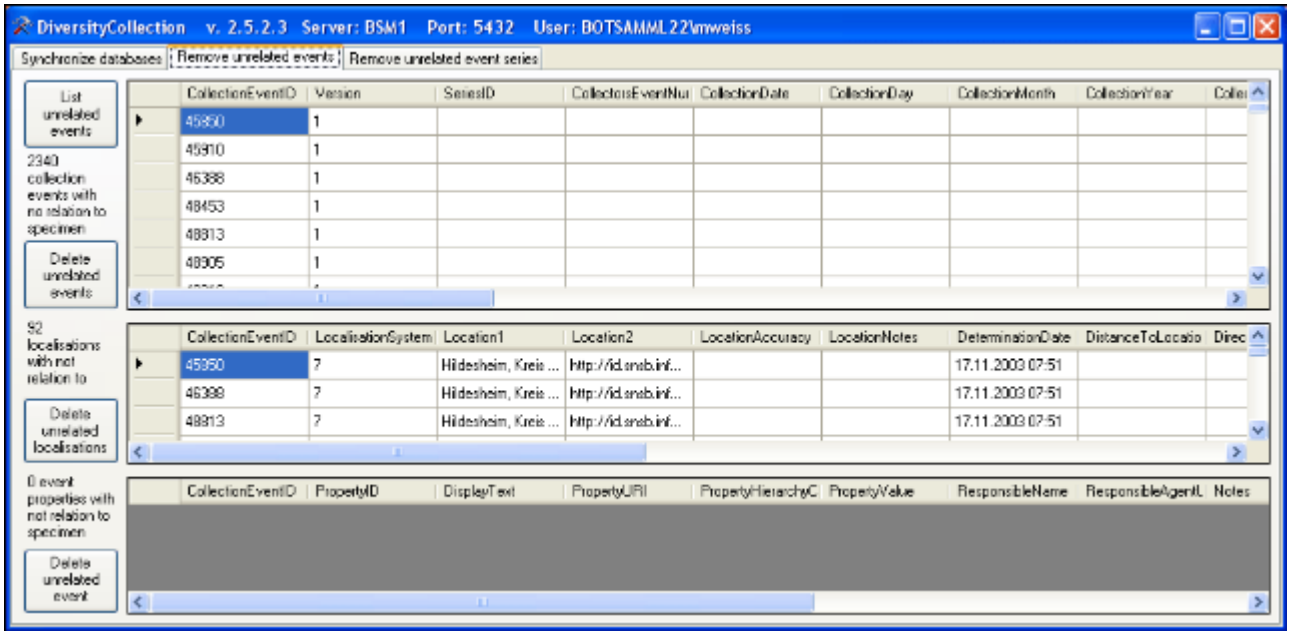
button. If you want to have a more detailed look on your data you have to check the **Include accession number** checkbox before starting the query. A button

will appear that will take you back to a single dataset in the database.

Maintenance - Unrelated data

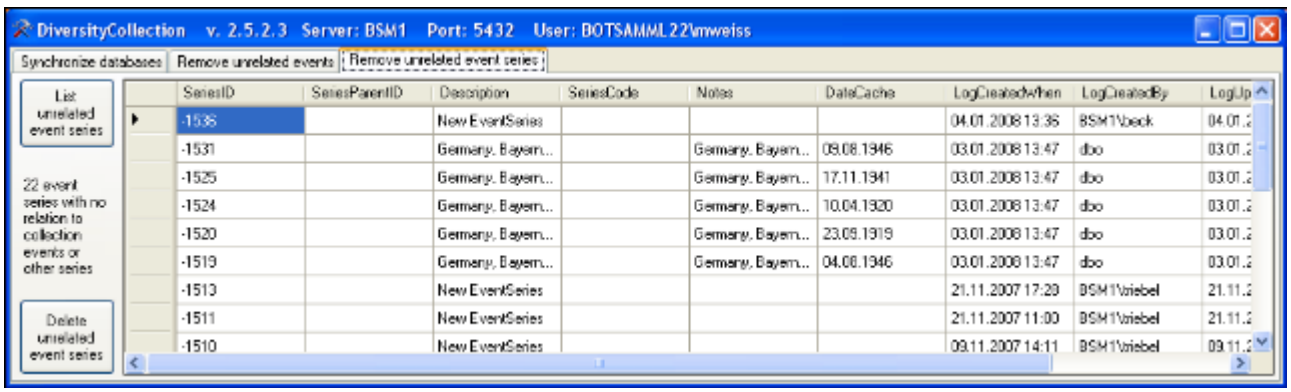
Collection events

Provided you have the proper rights, you can remove unlinked datasets in the tables CollectionEvent and CollectionEventSeries. To delete events that are not linked to any data in the database, use the tab page **[Remove unrelated events]**. These unrelated datasets may e.g. be derived from specimens that were transferred to another collection event. Click on the **[List unrelated events]** button to list all events that are not related to a specimen. The found collection events will be listed in the upper part of the form. The lower parts show the localisations and the event properties related to these collection events. These must be deleted first before you can delete the events related to these datasets. Otherwise only the events with no relations to localisations or properties will be deleted.



Collection event series

To delete collection event series with no relation to collection events or other event series, choose the tab page **[Remove unrelated event series]**. Click the **[List unrelated event series]** to list these series and the [Delete unrelated event series] button to delete them. .



Import and export

There are several import and export mechanisms:

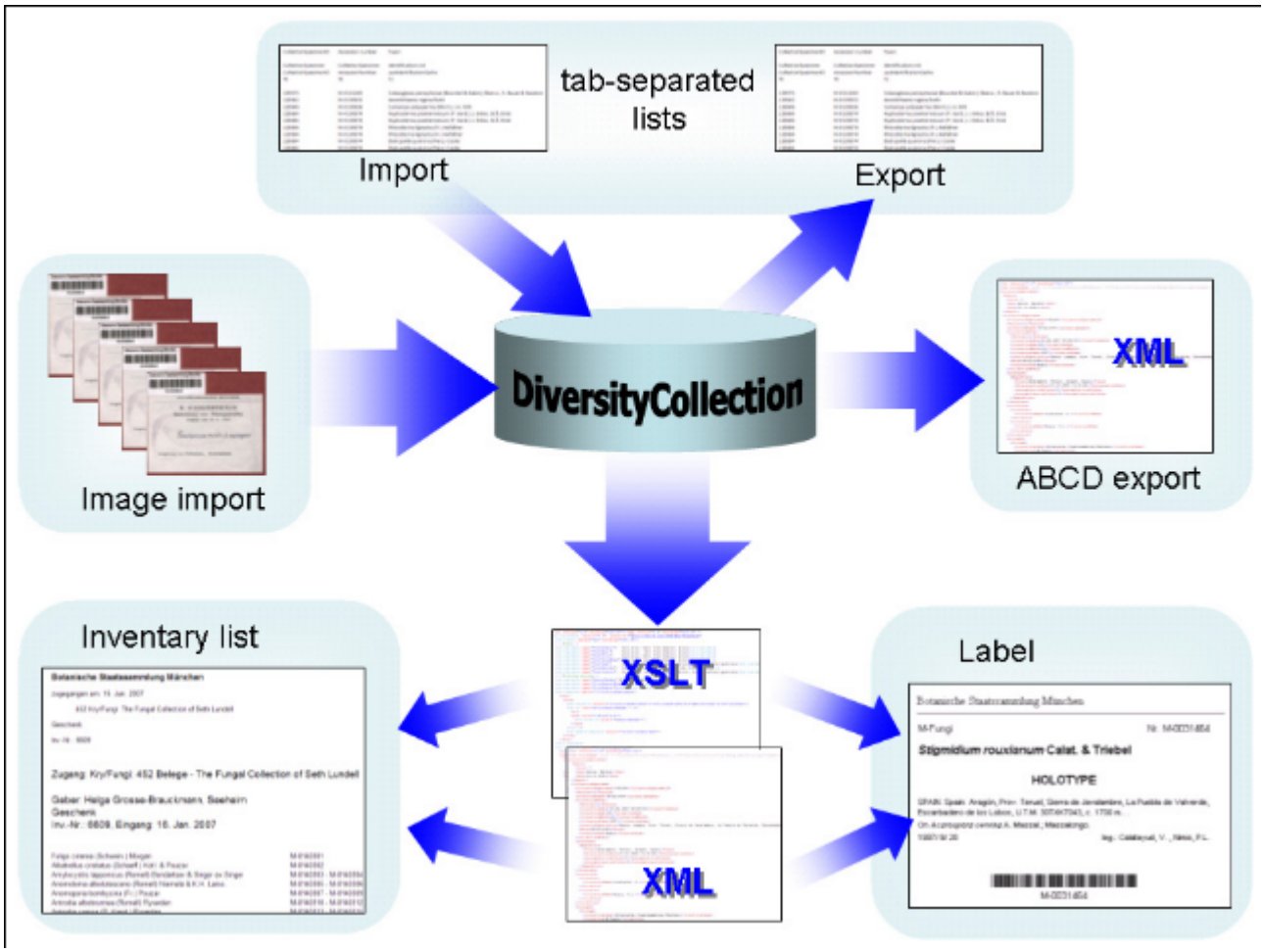
Import of specimen scans: Import image data where the accession number is a part of the file name can be imported together with default informations to the whole batch.

Import, export and reimport of tab-separated lists: Import data from foreign sources, export data and reimport after external editing.

Export of XML data according to the ABCD schema 2.06.

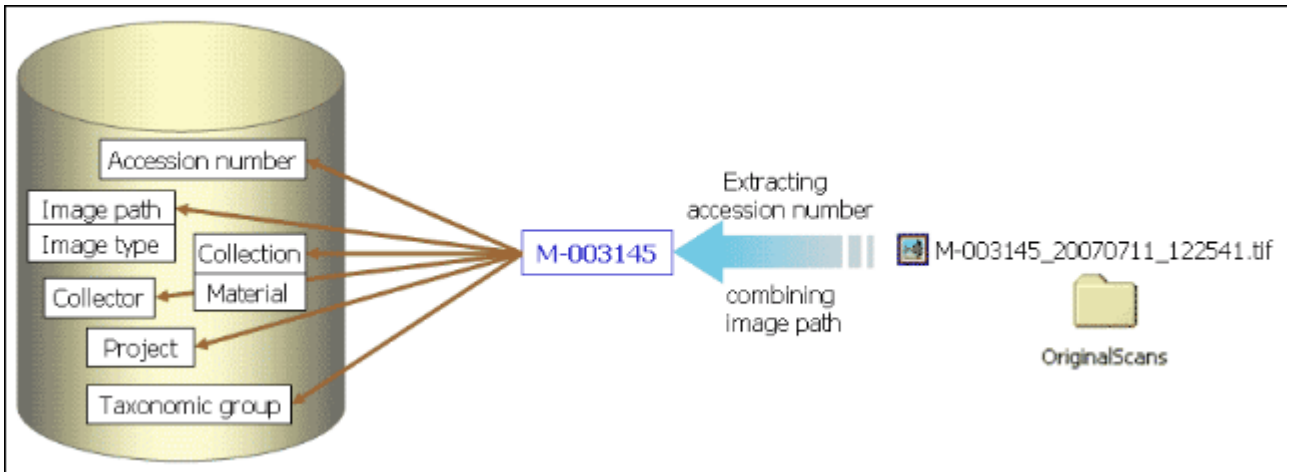
Generation of **labels**.

Generation of **inventary lists**.

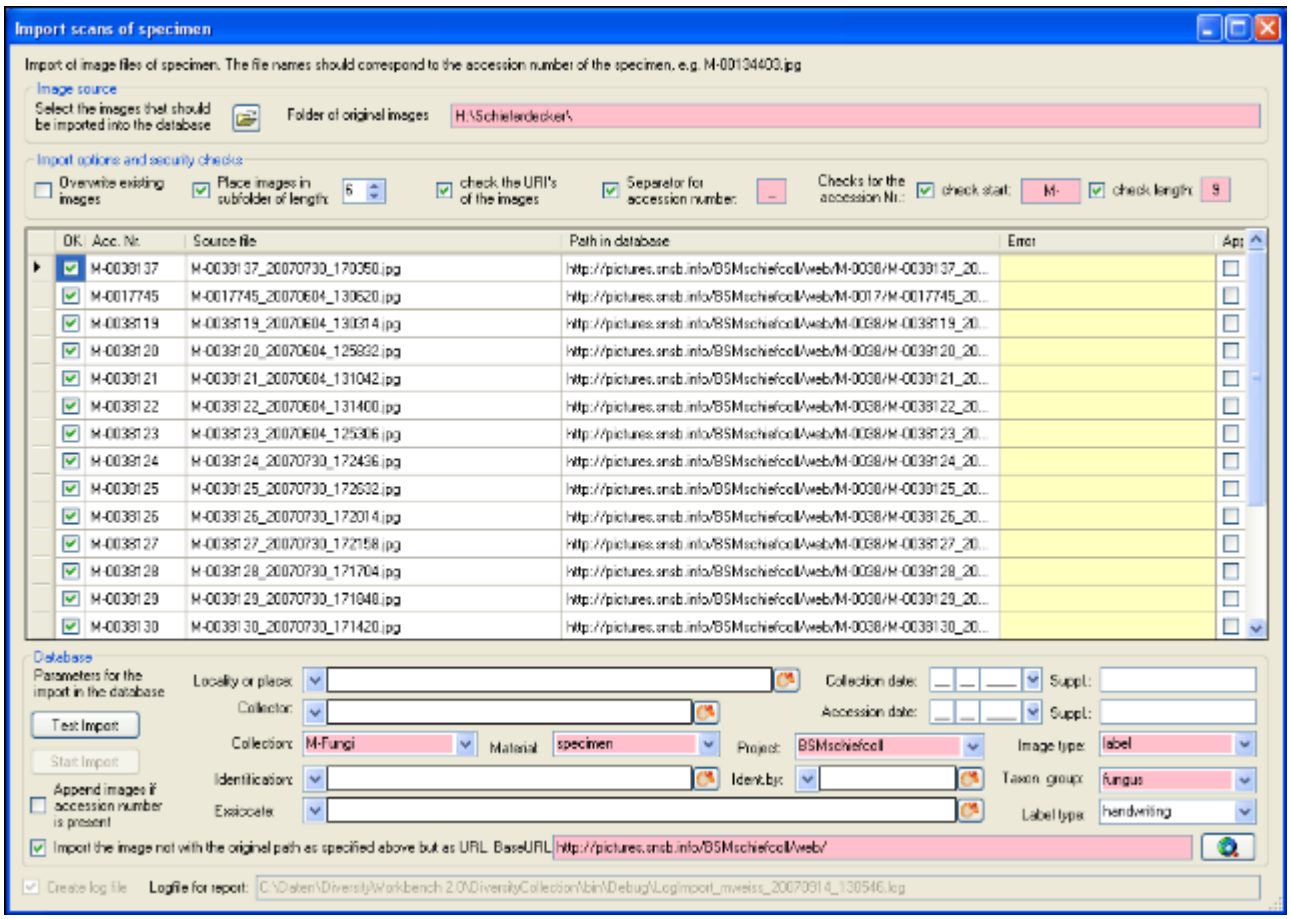


Import specimen scans

With this import routine, you can import new datasets along with new images into the database. To achieve this the image files should be named corresponding to the accession numbers of the datasets that should be imported (e.g. M-003145 in the example below) and an optional trailing identifier (e.g. _20070711_122541 in the example below). The image below shows an overview of the whole import process.



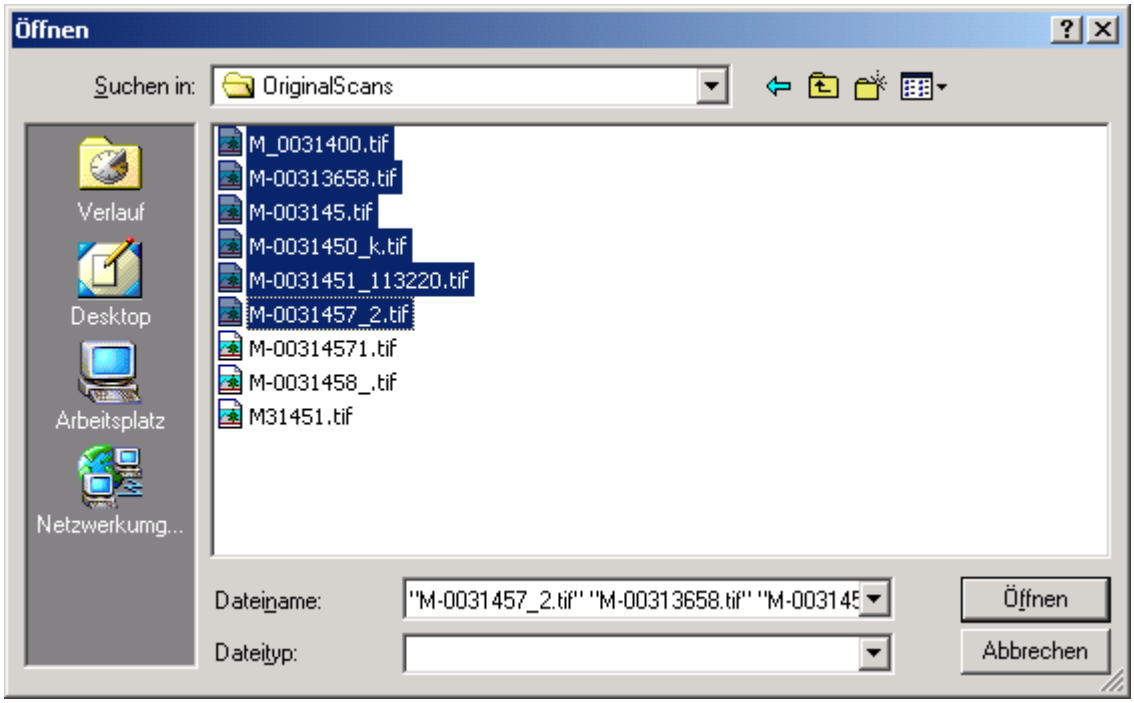
To import images like scanned labels from specimens choose **Data - Import - Specimen scans...** from the menu. A window for the import of images will be opened (see below).



The fields marked with red are mandatory.

Image source

To select the images you want to import in the database click on the button. A dialog will open, where you can select the images that you want to import into your database.



Select the files and close the window to enter the selection of the images in the list in the form.

Import options and security checks

If you want to replace datasets for images already in the database, check the **Overwrite existing images** checkbox. If the images are located in a subfolder, that is named according to the first characters of the accession number, check the **Place images in subfolder of length** checkbox and specify the length of the name of the folder. If you want to check the URIs of the images check the corresponding checkbox. The datasets will then only be imported if the images are available on the web. If the filename next to the accession number contains a trailing identifier, this must be separated by a unique character. Check the corresponding box and enter the separating character. If you want to check whether the accession numbers are correct you can check the start and the length of the accession number by checking the appropriate checkboxes.

Image list

The list shows the image files selected for the import. To test if all files satisfy the specified checks, click on the **Test import** button. If the file name passed all checks, the OK field for the image will be checked after the test. Otherwise an explanation for not passing the checks will be shown in the Error column.

The second column of the list shows the accession number extracted from the file name, the second column the file name of the original image file and the path written in the database. These paths may differ from the original if you select the option for setting a different **BaseURL**. If a dataset for an image already exists in the database and you selected the **Append images** option, the checkbox **Append** at the right end of the table will be checked.

Database

You can set several mandatory and optional values that will be written in the database for all

imported data. The fields marked with red are mandatory. These are the collection in which the specimens are located, the material categories of the specimens, the project, the content type of the scans (e.g. label) and the taxonomic group of the main organism in the specimens. If you want to refer to a web source for the labels instead of a local file you must give the base URL as well. Optional fields are a place according to the DiversityGazetteer and the date of the gathering event, the collector, the accession date, an identification, an exsiccatal series and the type of the label. For explanation of the buttons see module related entries .

To test the import and whether the images are present in the database click on the

button. If everything is fine click on the

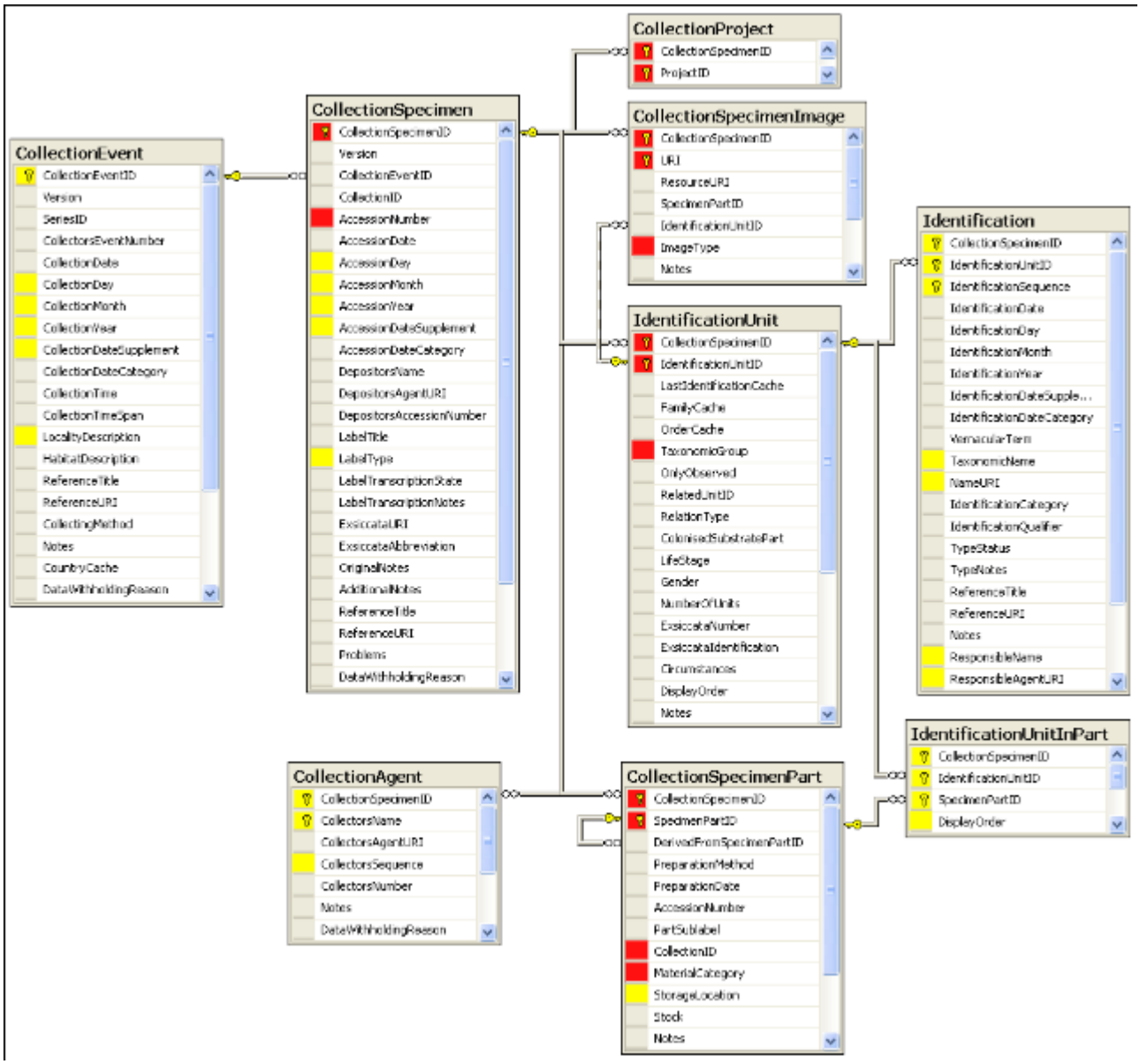
button to start the import.

Logfile: To log the list of imported images and any errors during the import, check the Create log file checkbox. This will create a log file with your name, the date and time of the import in the directory where the image files are located.

The program will check, whether an accession number is already present in the database. It will only import the data, if the option **Append images if accession number is present** is checked. Otherwise these data will not be imported.

If the images will be provided by a webserver, check the appropriate checkbox and specify the folder or click on the button of search for the website.

In tIn the overview below the mandatory fields are marked with red, the optional fields with yellow.


CollectionProject
CollectionSpecimenID
ProjectID
CollectionSpecimen
CollectionSpecimenID
Version
CollectionEventID
CollectionID
AccessionNumber
AccessionDate
AccessionDay
AccessionMonth
AccessionYear
AccessionDateSupplement
AccessionDateCategory
DepositorsName
DepositorsAgentURI
DepositorsAccessionNumber
LabelTitle
LabelType
LabelTranscriptionState
LabelTranscriptionNotes
ExsiccataURI
ExsiccataAbbreviation
OriginalNotes
AdditionalNotes
ReferenceTitle
ReferenceURI
Problems
DataWithholdingReason
CollectionSpecimenImage
CollectionSpecimenID
URI
ResourceURI
SpecimenPartID
IdentificationUnitID
ImageType
Notes
CollectionEvent
CollectionEventID
Version
SeriesID
CollectorsEventNumber
CollectionDate
CollectionDay
CollectionMonth
CollectionYear
CollectionDateSupplement
CollectionDateCategory
CollectionTime
CollectionTimeSpan
LocalityDescription
HabitatDescription
ReferenceTitle
ReferenceURI
CollectingMethod
Notes
CountryCache
DataWithholdingReason
IdentificationUnit
CollectionSpecimenID
IdentificationUnitID
LastIdentificationCache
FamilyCache
OrderCache
TaxonomicGroup
OnlyObserved
RelatedUnitID
RelationType
ColonisedSubstratePart
LifeStage
Gender
NumberOfUnits
ExsiccataNumber
ExsiccataIdentification
Circumstances
DisplayOrder
Notes
Identification
CollectionSpecimenID
IdentificationUnitID
IdentificationSequence
IdentificationDate
IdentificationDay
IdentificationMonth
IdentificationYear
IdentificationDateSupple...
IdentificationDateCategory
VernacularTerm
TaxonomicName
NameURI
IdentificationCategory
IdentificationQualifier
TypeStatus
TypeNotes
ReferenceTitle
ReferenceURI
Notes
ResponsibleName
ResponsibleAgentURI
IdentificationUnitInPart
CollectionSpecimenID
IdentificationUnitID
SpecimenPartID
DisplayOrder
CollectionAgent
CollectionSpecimenID
CollectorsName
CollectorsAgentURI
CollectorsSequence
CollectorsNumber
Notes
DataWithholdingReason
CollectionSpecimenPart
CollectionSpecimenID
SpecimenPartID
DerivedFromSpecimenPartID
PreparationMethod
PreparationDate
AccessionNumber
PartSublabel
CollectionID
MaterialCategory
StorageLocation
Stock
Notes

Import tab-separated lists

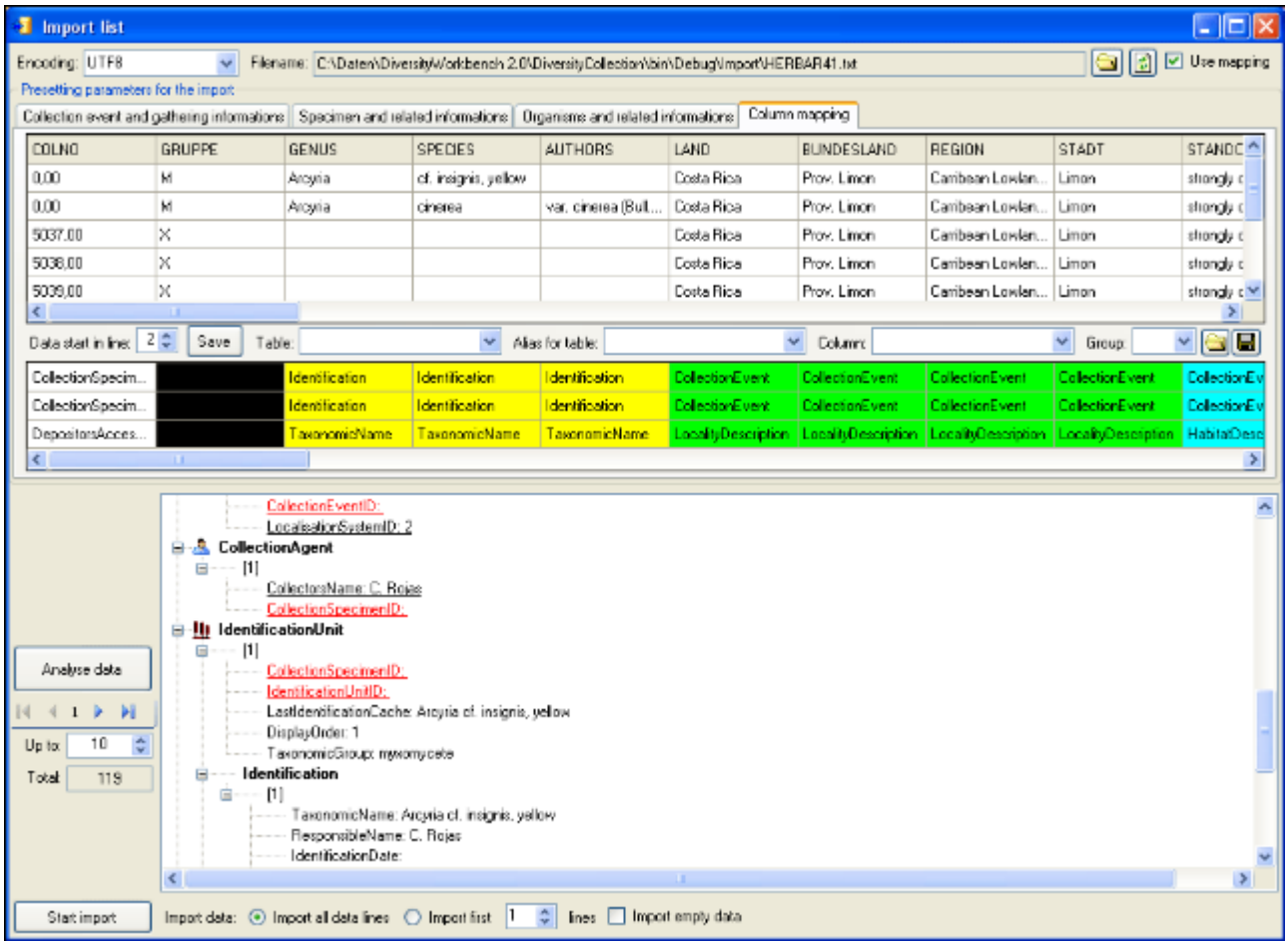
With this import routine, you can import data as tab-separated lists into the database. Choose **Data -> Import -> Import list...** from the menu to open the window for the import. In the window click on the button to select the file with the data you want to import. Use the proper **[encoding]** to ensure that all special characters will be transferred correctly. If you change the encoding after opening the file, click on the button to reload the file with the new encoding.

The content of the file will be shown in the upper part of the **[Column mapping]** tab page. Use the **[Data start in line]** to set the area for the import. Preceeding lines will be ignored and depicted with a gray background as shown below. Every column in your file must be either ignored of mapped to a column in the database. The lower part of the mapping section shows your mapping, while the upper part contains the first lines of your file. You can either use a prepared column mapping or create a new one. To import a previous column mapping, click on the button and choose one of the XML-files.

Column mapping

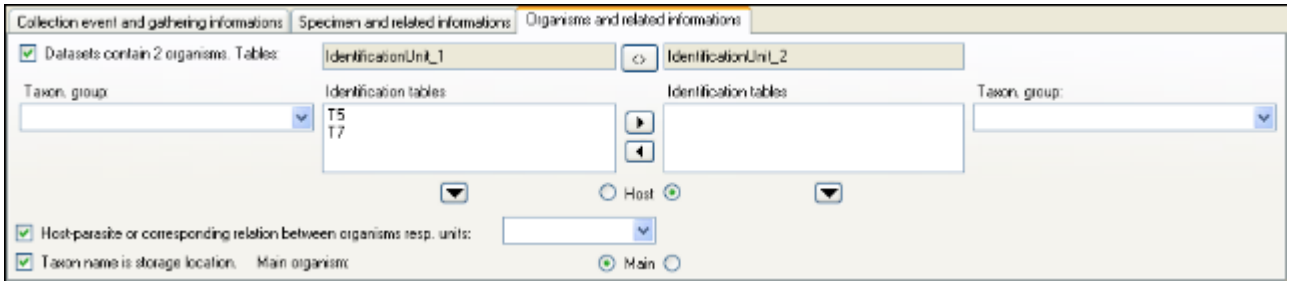
For a column that should not be imported, choose the black color from the group as shown for the second column in the image below. These columns will be ignored for the analysis and the import. For all other columns choose the **[Table]** and the **[Column]** your data correspond to. If you want to import two different datasets in the same table, you have to change the **[Alias for table]** to a unique value for this dataset.

If some columns should be transferred into one field, you have to choose a group for these column as shown below for columns 3 to 5, 6 to 9, ... To do this, just choose a color from the group combobox other than white or black. To save the current mapping, click on the button.



To analyse the data in the file click on the **[Analyse]** button. During the analysis the program may ask you to give additional information, like the taxonomic groups of the imported organisms. In the upper part of the window you can add informations that should be imported together with your data from the file. The available options depend on the structure of your data.

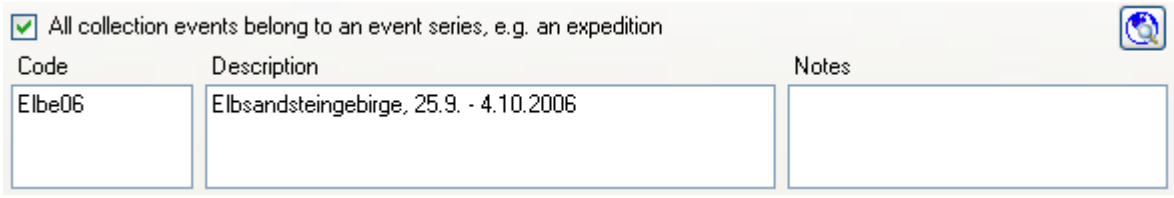
The image below shows the tab page where you can define the relations between two organisms. This might be necessary if e.g. your data contain informations of parasites and hosts. The data tables that are related to an identification are listed with their aliases in the lists for the identification tables. Use the ▶ and ◀ buttons to move them between the lists. Use the ▼ buttons to change the sequence within a list. With the **[Host]** option ○ Host ◉ you define the host and the **[Main]** option defines which organism will be chosen as the first to be printed e.g. on a label.



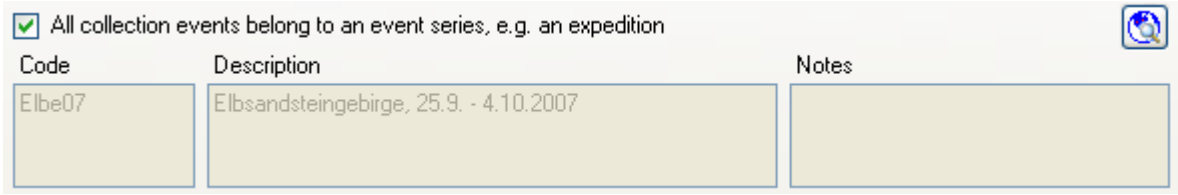
By default one collection event will be created for each dataset.

If the events should be joined following the entries in the file, you can choose the second option as shown below. The import will create a new event if there is any change in the data related to the collection event, like the description of the locality, the altitude, the coordinates etc.

If all the dataset are belonging to one event series, e.g. to one expedition, you can choose the option as shown below and enter the code and description of the event series in the corresponding fields.



If you want to use an existing event series, click on the button and select a series from the form. The code and description of the selected series will be shown as in the image below.

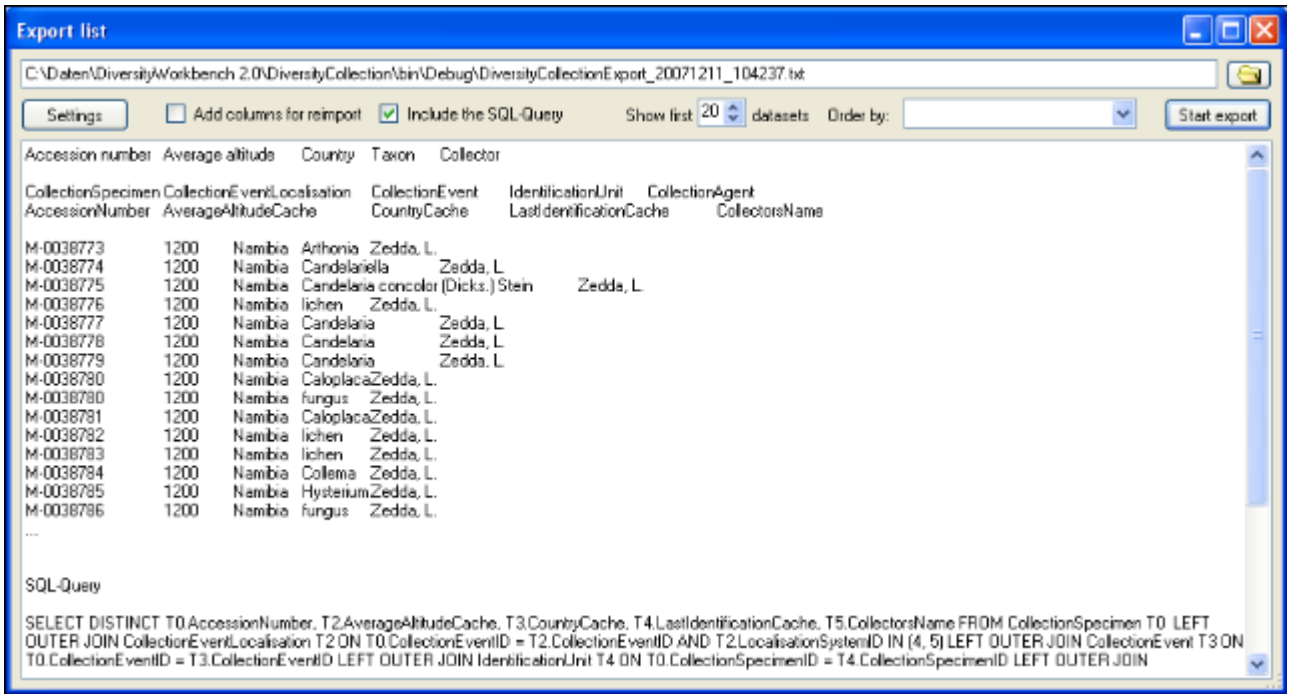


After your data where successfully analysed, the window will show you the result as shown in the image below. During the analysis, the program will add missing tables and columns necessary for the import. The first dataset will be shown in the form. If you want to check more datasets, set the number of the last position you want to check and click on the **[Analyse]** button. The data are arranged according to the tables in the database in which they will be imported. The underlined fields belong the the primary keys of the respective tables and are red if missing. These missing values will be generated during the import. If a dataset contains no values and will therefore not be imported into the database, the colums will be shown in blue. To import these entries in any case, choose the **[import empty values]** option. To browse through the datasets in the preview use the 2 buttons.

To import the data, click on the **[Start import]** button. If you want to import just a part of the data, e.g. for a test, choose the **[Import first ... lines]** option.

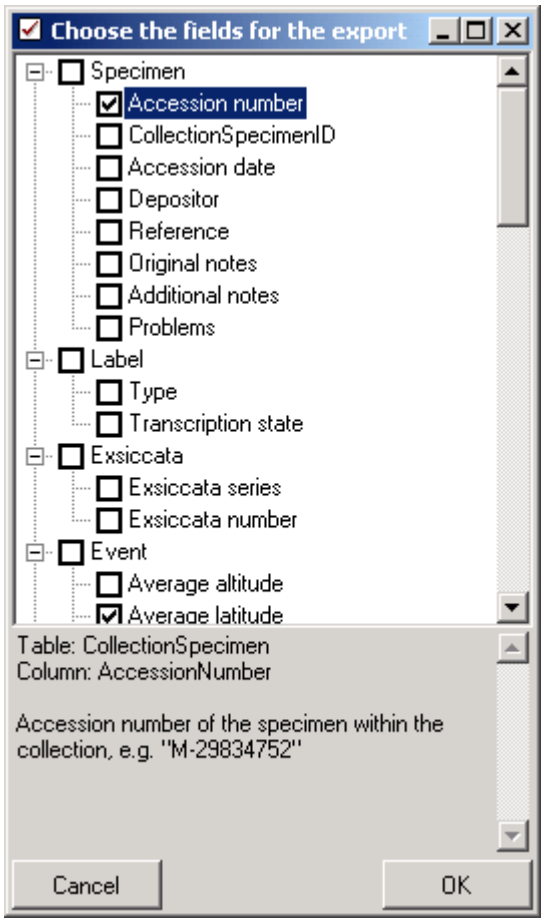
Export as tab-separated lists

You can export the content of the datasets listed in the specimen list into a tab-separated list. Choose **Data - Export - Export list...** from the menu. A window as shown below will open. To change the preset path of the export file use the button.



To start the export click on the **[Start export]** button. A file will be created in your application directory, containing the exported data. A preview of the data is shown in the lower part of the form. If you check the **include columns for reimport** option, the header will contain an additional line for table names which you can use for reimporting the data. If you check the **include the SQL-Query** option, the Text of the command for selecting the datasets will be attached at the end of your report. The form will show the first lines as set in the Show first ... datasets (range: 1 - 99) as a preview. During the first export, the list for the fields for sorting the results **[Order by:]** will be filled. So if you need sorted results just restart the export after selecting the field after the first export.

To choose the fields you want to see in the export click on the Settings button. A window will open as shown below, where you can choose the fields you want to export.



Please keep in mind, that only the datasets listed in the **specimen list** of the main form will be exported. If for example you set the maximal number for the specimen list to 100, only these datasets will be exported, even if the number of datasets related to your query will be more than 100.

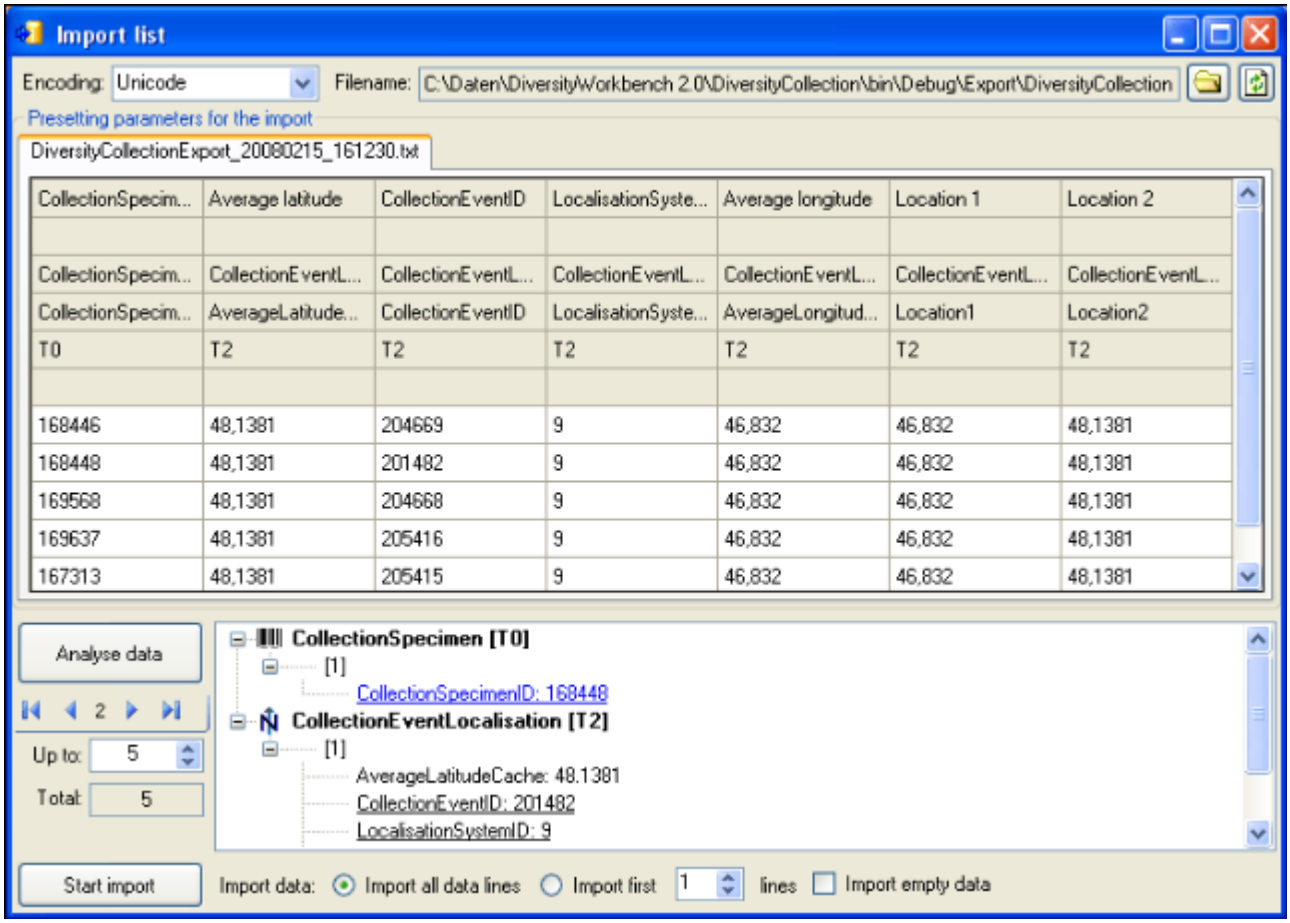
Please be aware, that you will get "**cross joins**" between the data in your database as in the example shown below. If for example you start an export containing taxa and collectors and you have 3 taxa and 2 collectors in a dataset this will result in 6 lines in the export: every taxon combined with every collector (3 x 2 = 6). To evaluate the data you have to group the results accordingly. The first line in the file corresponds to the description as shown in the form above for the selection of the export fields. The second line shows the tables and the third line the column within the database. The following lines contain the data. If you want to avoid these cross joins and need only one line for each dataset, please use the export funktion of the **grid mode**.

Accession number	**Taxon**
CollectionSpecimen	**IdentificationUnit**
AccessionNumber	**LastIdentificationCache**
M-0013579	Arthrocladiella mougeotii (Lev.) Vassilkov
M-0013579	Arthrocladiella mougeotii (Lev.) Vassilkov
M-0013579	Blumeria graminis (DC.) Speer
M-0013579	Blumeria graminis (DC.) Speer
M-0013579	Lycium barbarum
M-0013579	Lycium barbarum

Reimport tab-separated lists

With this import routine, you can reimport data in tab-separated lists that had been exported from the database. Choose **Data -> Import -> Reimport list...** from the menu to open the window for the reimport. To reimport data, these must have been exported using the option **[add columns for reimport]** to ensure correct header lines in your file. As the client provides no possibility to change data on the basis of lists, you may export the respective data to a tab-separated list, change your values and reimport the data.

In the window click on the button to select the file with the data you want to import. Use the proper **[encoding]** to ensure that all special characters will be transferred correctly. If you change the encoding after opening the file, click on the button to reload the file with the new encoding. The content of the file will be visible as shown below. For a reimport, the data will always start in line 7, indicated by a gray background of the header lines.



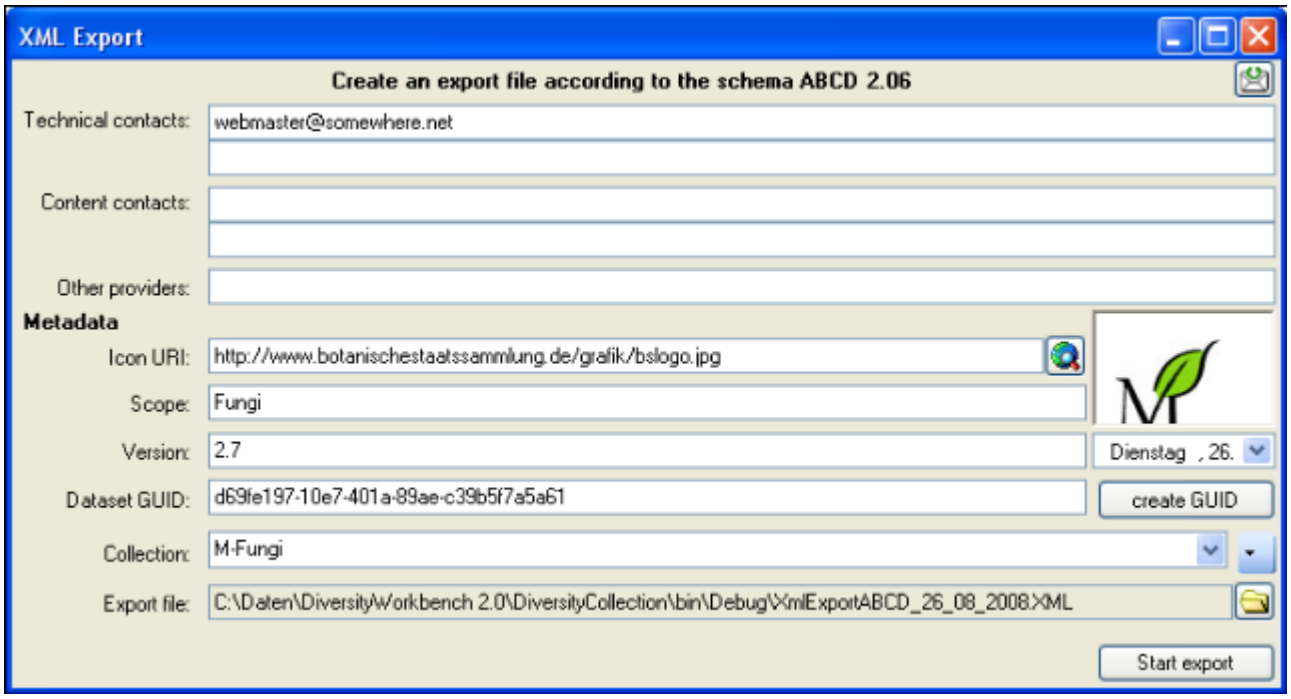
To analyse the data in the file click on the **[Analyse]** button. After a successful analysis, the window will show you the result as shown in the image below. The first dataset will be shown in the form. If you want to check more datasets, set the number of the last position you want to check and click on the **[Analyse]** button. The data are presented according to the tables in the database where they will be inserted. Underlined columns belong to the primary key of the respective tables and are red if missing. These missing values will be generated during the import. If a dataset contains no values and will therefore not be imported into the database respectively updated, the colums will be shown in blue. To import these entries in any case, choose the **[import empty values]** option. To browse through the datasets in the preview use the buttons.

To import your data, click on the **[Start import]** button. If you want to import just a part of the data, e.g. for a test, choose the **[Import first ... lines]** option.

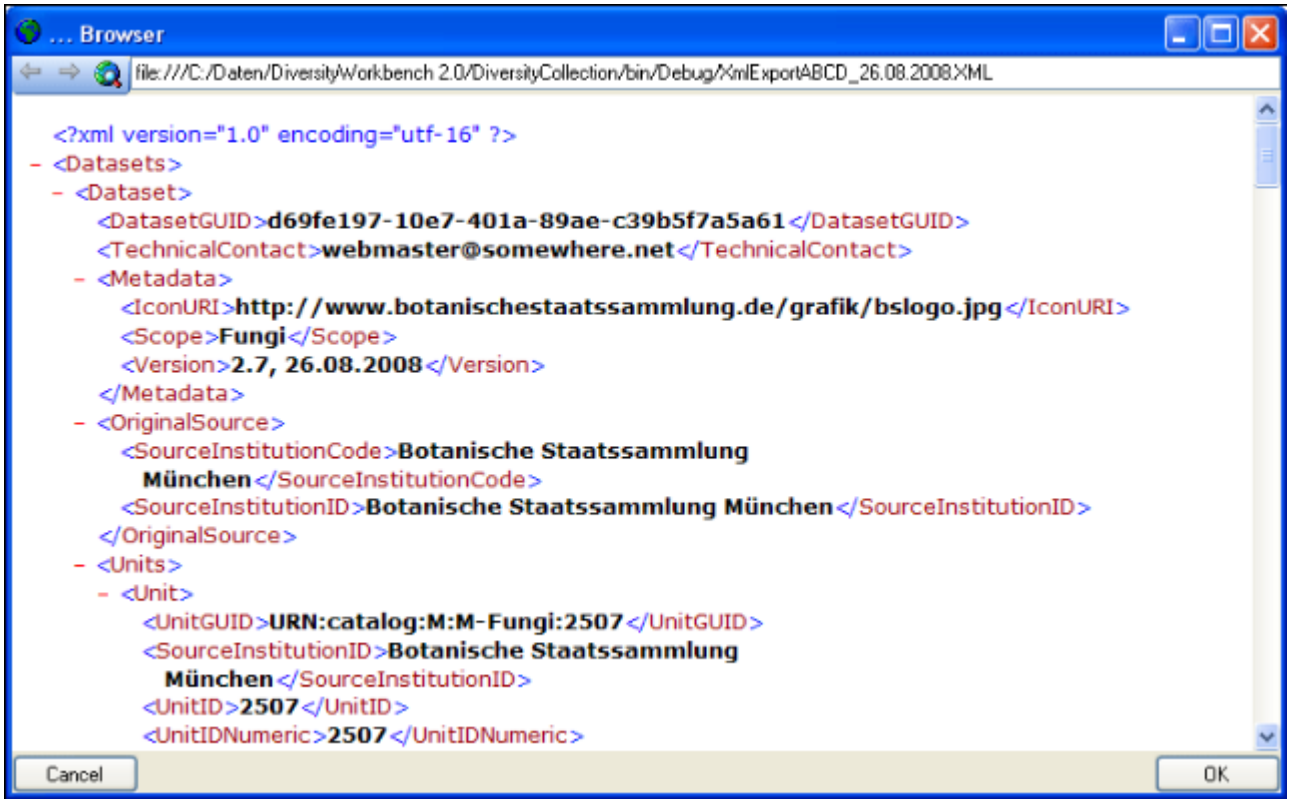
Export ABCD

In the current version only the main data from DiversityCollection will be exported to ABCD.

To export the data of the specimen selected in the specimen list following the ABCD schema 2.06 choose **Data -> Export -> XML (ABCD Schema)...** from the menu. A window as shown below will open where you can set some additional parameters defined in ABCD resp. BioCASE.

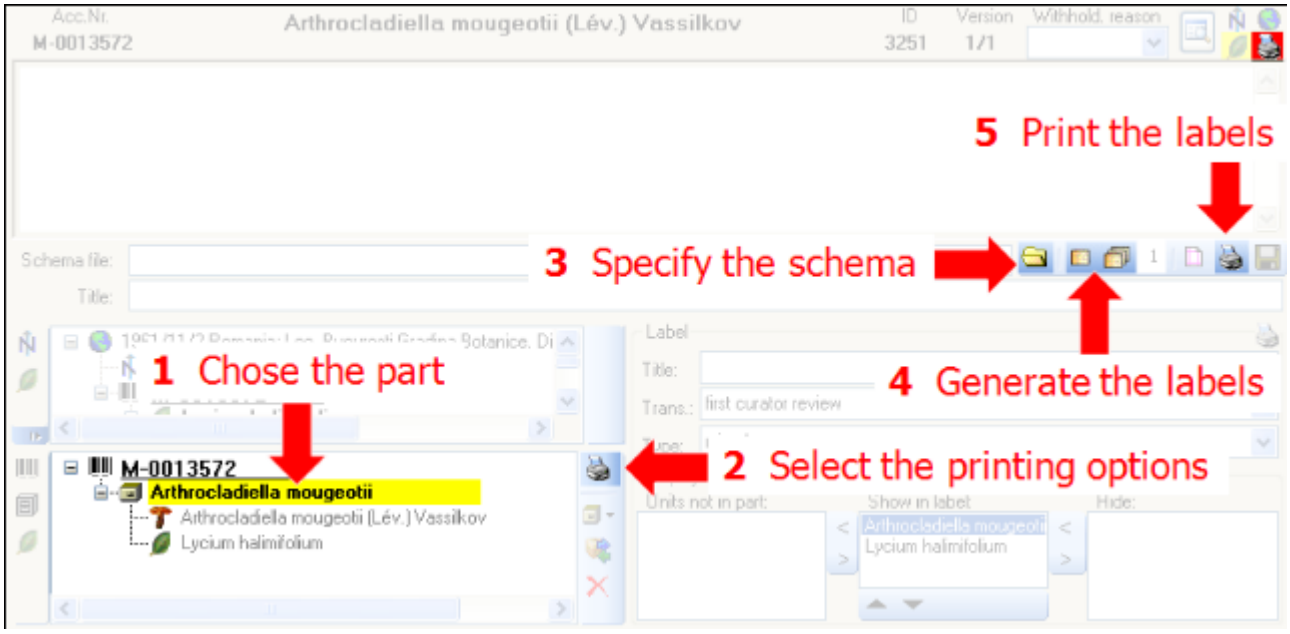


To start the export click on the **Start export** button. The data will be exported into a file in your application directory. Click on the button to inspect the exported data (see below).

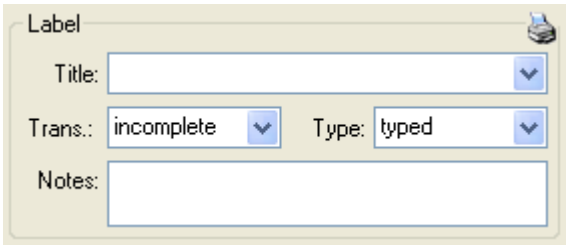


Label

If there is only one part in your specimen click on the printer symbol in the button to switch to the printing mode. If your specimen contains more then one part, choose the part of the specimen for which the label should be generated and click on the button in the right panel. The image area will switch to the label view and in the data area the details related to the label will be accessible. The sequence is shown in the image below.



Additional information about a label are entered in the label section (see image below). The data are stored in the table CollectionSpecimen.



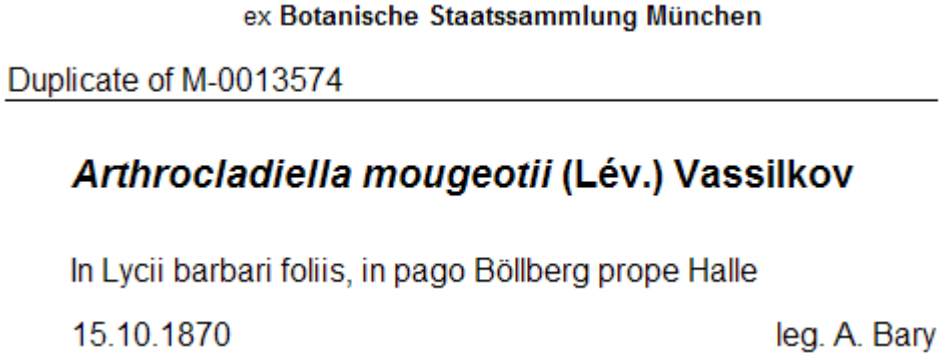
The organisms of a specimen are printed on a label according to the display order.

To print a label for a specimen you have to select a schema file. There are default schema files available in the folder **LabelPrinting/Schemas** in your application directory. Click on the button to open the directory. You will find several prepared schema files among which you can choose resp. change them to your own needs or create new ones. The schema file **LabelTemplates.xslt** provides templates for the other schema files. You may give a title for the print in the field Title. From the Collection and MaterialCategory available for the selected specimen choose one from the list (**Coll./Mat.**). To generate the label for the current specimen click on the button. To generate labels for all specimens in your query click on the button. If you need duplicates of your labels change the number in the duplicates box 3 to the desired value. You can print 1 - 99 duplicates of one label. If there are more than 20 specimens in the list, you get a warning whether you really want to create all these labels, as this could be somewhat time consuming. The labels are generated as XML files with

XSLT-schema files, transformed to HTML-files and depicted in a browser. To print the label click on the button.

If you want to print labels for duplicates that are stored in a different collection, the duplicate should be a child of the original specimen as shown in the example below.

IDepending on the schema you use, the label will be marked as duplicat and contain a reference to the original specimen (see below).



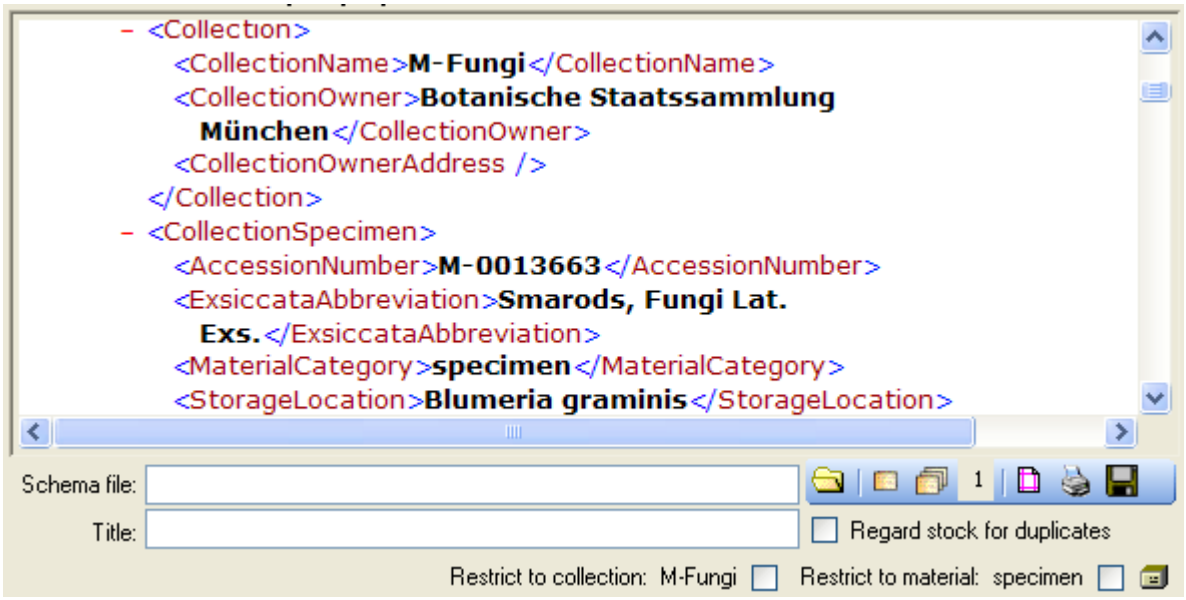
If you want to save the generated files for later printing click on the button to do this.

Note that the program will by default create a file **Label.xml** and in case a schema file is specified a file **Label.htm** in the **LabelPrinting** directory that will be overwritten everytime you generate a new label. So you have to save the file under a different name or in a different directory to prevent the program to erase these data.

If you use **Code 39** for your labels and want to print the barecodes on the labels you need the font code39.ttf , which is included in the DiversityCollection packet. Place this font in the folder where your fonts are stored (e.g.: C:\WINNT\Fonts). If the font is not available, the Barcode will appear as the accession number between two '*' signs.

If you want to print labels for all the specimens in the specimen list you can restrict these to the collection and the material category of the current specimen part (see image above).

If you do not select a schema file, i.e. the textbox **Schema file:** is empty, you will see the generated XML-file as shown in the image below. The XML file is the base for all label types you want to generate. To create your own labels just design your own XSLT-schema file. See e.g. http://www.w3.org/TR/xslt for further informations about schema files.



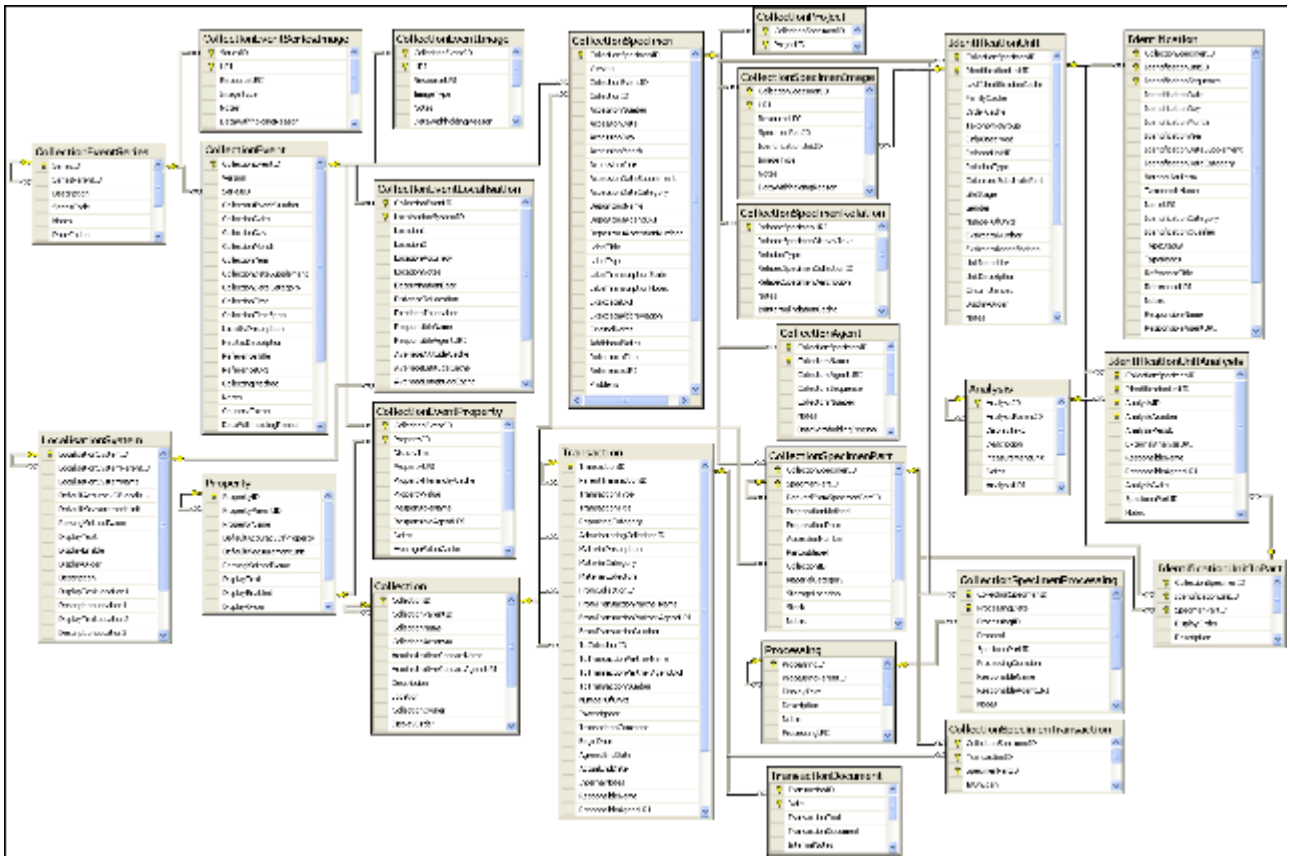
Database

The database for DiversityCollection is based on Microsoft SQL-Server 2005.

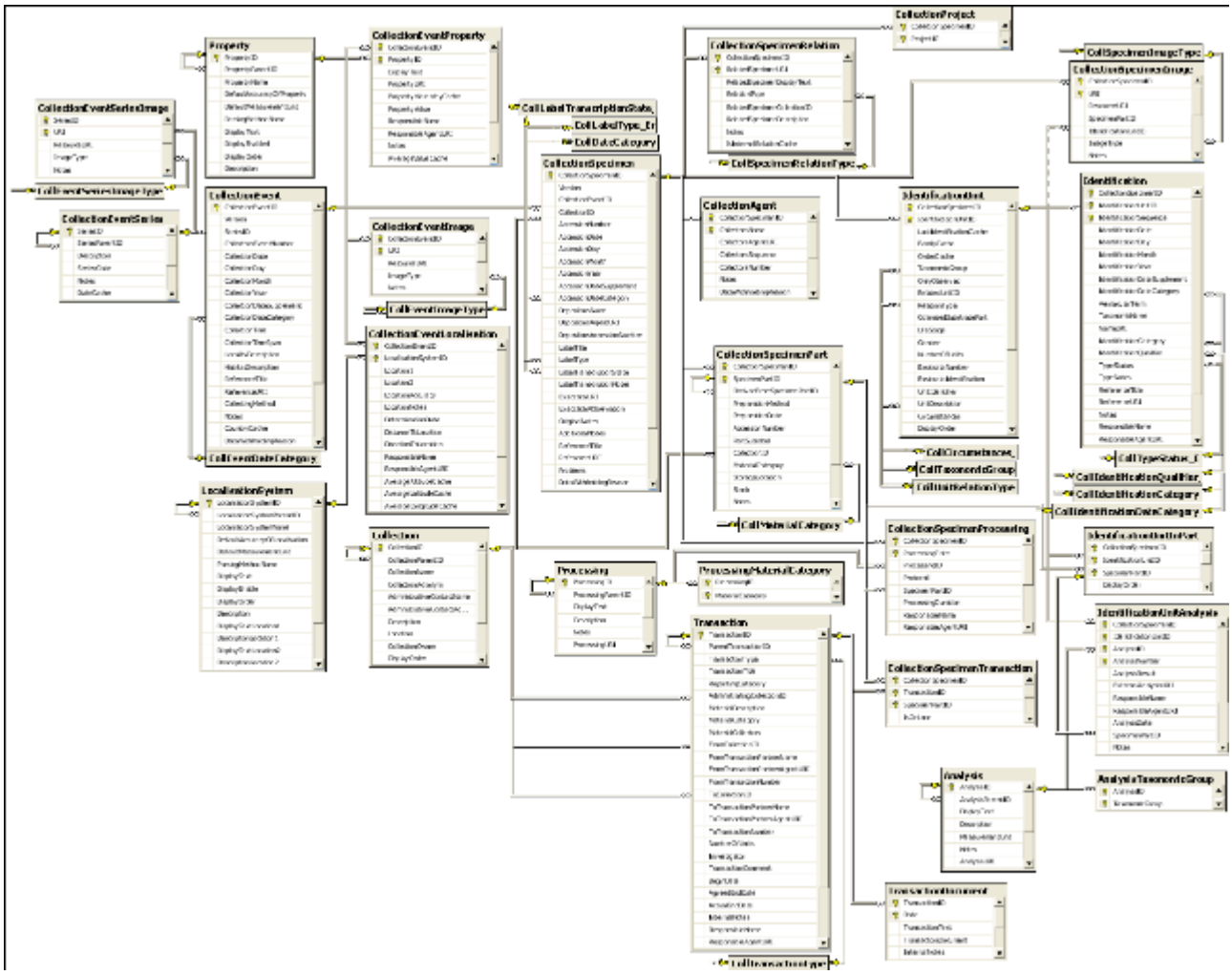
Organisation of the data

The main tables of the database are CollectionEvent corresponding to the event of the collection and CollectionSpecimen holding the specimens collected. Connected to these tables you find tables for additional informations.

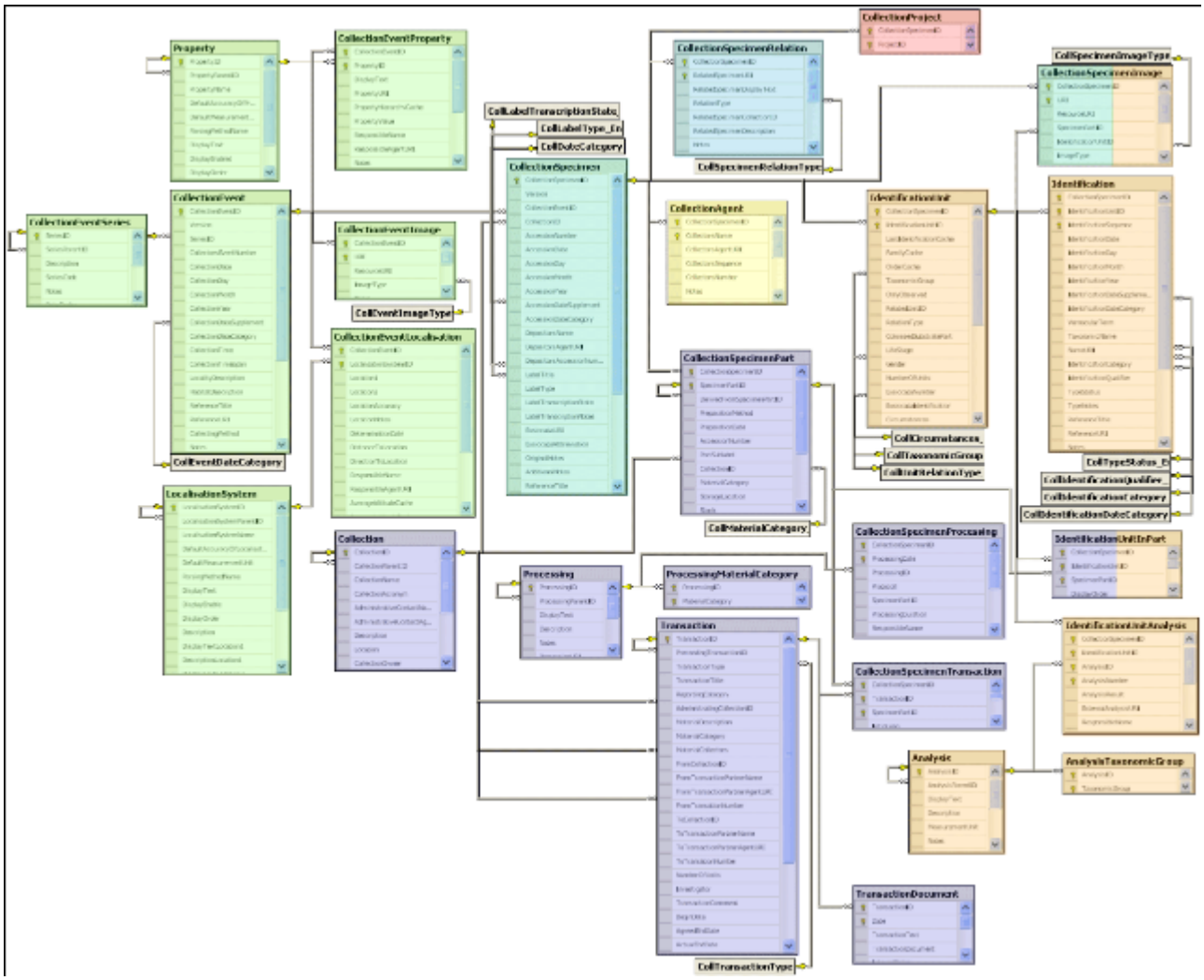
The image below shows the main tables of the database.



The structure of the whole database is shown in the image below.



In the graphic below the tables of the database are marked according to their logical groups. The central is the table CollectionSpecimen holding the data of the specimen like e.g. the accession number. In the left part you find the tables containing data related to the collection event, and in the right part tables related to the organisms and their identifications. Further logical groups are relation, collectors, projects and storage.



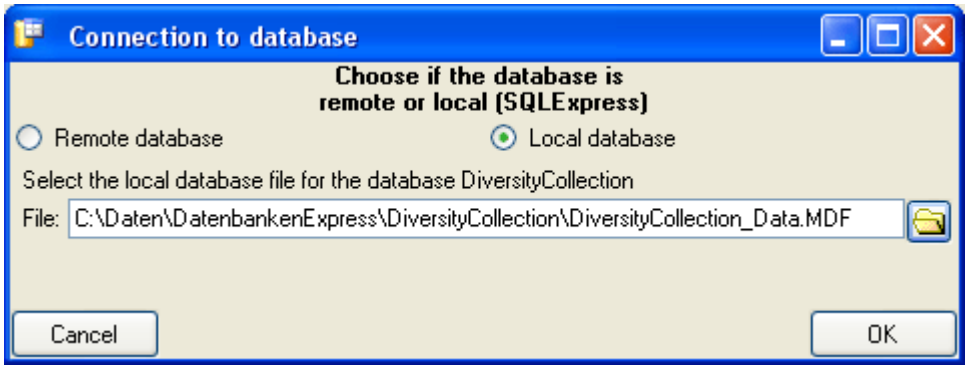
Further details: [tables](#) , [application tables](#), [access](#).

Database - access

The database engine for DiversityCollection is Microsoft SQL-Server 2005.

If you are connected to a database this is indicated by the icon of the connection button in left upper corner. If you are not connected this will be indicated by the icon . To access any database, you must specify the server where the database is located. For the configuration of this connection choose **Connection**, **Database...** from the menu or click on the button.

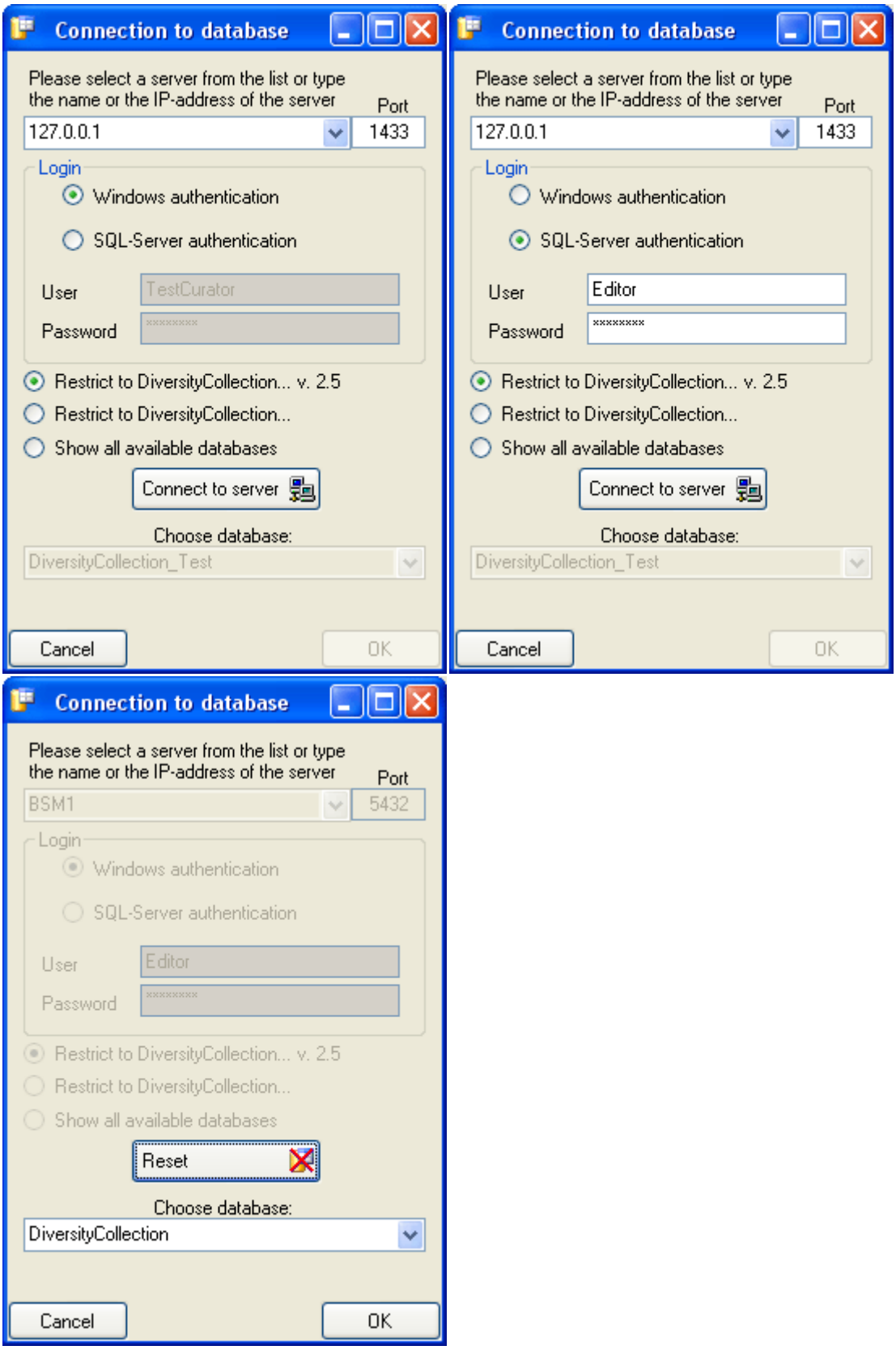
If you want to use a database on your local computer (SQL Server Express), start the program DiversityCollection.exe and in the main window click on the button or choose **Connection -> Database ...** from the menu. A window as shown below will open. Here choose the option **Local database** and select the database file in your directory.



If you want to use a database on a central server or remote computer choose the option **Remote database** and set the connection parameters as described below.

Database name, IP-address and Port

A dialog will open, to specify the name or IP-address and port number of the server and to select the authentication mode. You can either choose Windows authentication (see left image below) or SQL-Server authentication (see below - central middle image).



The standard port number for SQL-Server is 1433 and will be set as a default. If the database server is configured using a port different from that port, you must give the port number in the field Port. Click on the

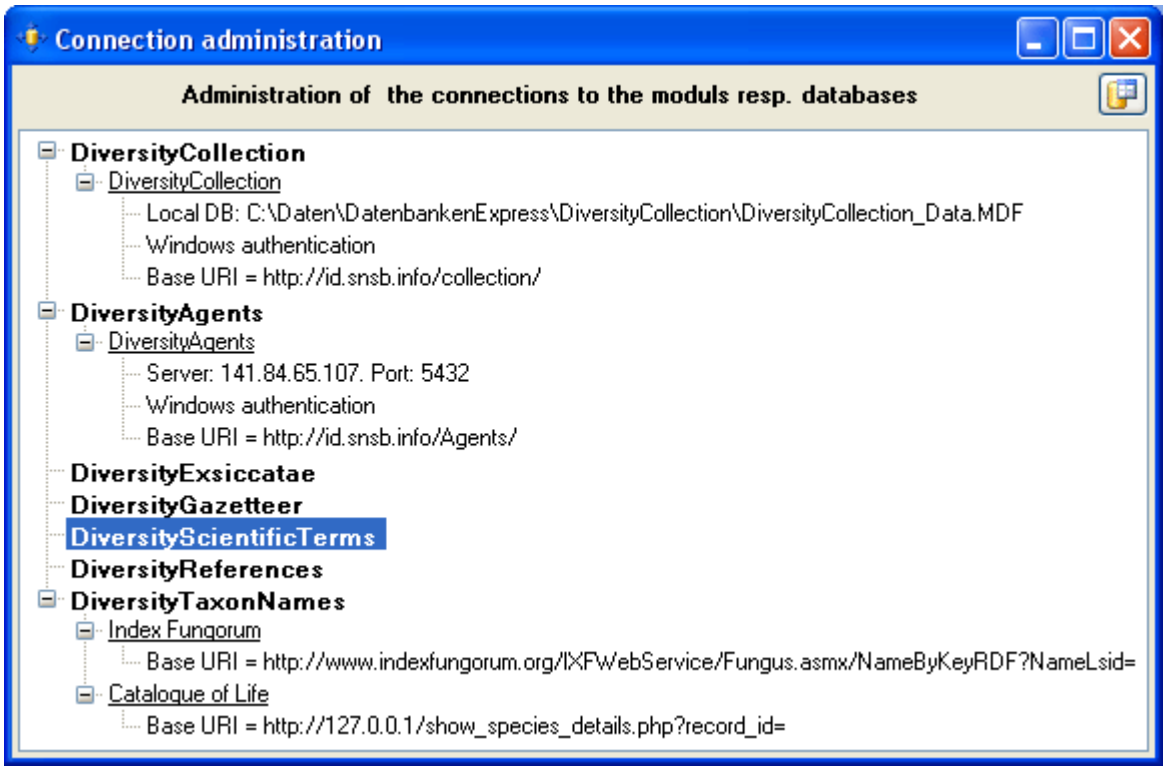
button to connect to the server. If the connection informations are valid, you can choose a database from the server from the combobox at the base of the window (see right image above). To restart the connecting

process click on the

button.

Module connections

The program will automatically try to get connection to all the modules within the Diversity Workbench. To edit these connections choose Connection - Module connections ... from the menu. A form as shown below will open, where you can edit these connections.



To edit a connection, select it in the tree and click on the button.

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Table Analysis

Analysis types used within the database

Column	Data type	Description
AnalysisID	int	ID of the analysis (Primary key)
AnalysisParentID	int	Analysis ID of the parent analysis if it belongs to a certain type documented in this table
DisplayText	nvarchar (50)	Name of the analysis as e.g. shown in user interface
Description	nvarchar (MAX)	Description of the analysis
MeasurementUnit	nvarchar (50)	The measurement unit used for the analysis, e.g. mm, µmol, kg
Notes	nvarchar (MAX)	Notes concerning this analysis
AnalysisURI	varchar (255)	URI referring to an external documentation of the analysis
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table AnalysisTaxonomicGroup

The types of analysis that are available for a taxonomic group

Column	Data type	Description
AnalysisID	int	Analysis ID, foreign key of table Analysis.
TaxonomicGroup	nvarchar (50)	Taxonomic group the organism identified by this unit belongs to. Groups listed in table CollTaxonomicGroup_Enum (= foreign key)
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table Collection

The collections where the specimen are stored

Column	Data type	Description
CollectionID	int	Unique reference ID for the collection (= Primary key)
CollectionParentID	int	For a subcollection within another collection: CollectionID of the collection to which the subcollection belongs. Empty for an independent collection
CollectionName	nvarchar (255)	Name of the collection (e. g. 'Herbarium Kew') or subcollection (e. g. 'cone collection', 'alcohol preservations'). This text should be kept relatively short, use Description for additional information
CollectionAcronym	nvarchar (10)	A unique code for the Collection, e.g. the herbarium code from Index Herbariorum
AdministrativeContactName	nvarchar (500)	The name of the person or organisation responsible for this collection
AdministrativeContactAgentURI	varchar (255)	The URI of the person or organisation responsible for the Collection e.g. as provided by the module DiversityAgents
Description	nvarchar (MAX)	A short description of the collection
Location	nvarchar (255)	Optionally location of the collection, e.g. the number within a file system or a description of the room(s) housing the (sub)collection

CollectionOwner	nvarchar (255)	The owner of the collection as e.g. printed on a label, should be given if CollectionParentID is null
DisplayOrder	smallint	The order in which the entries are displayed. The order may be changed at any time, but all values must be unique.
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionAgent

The collector(s) of collection specimens

Column	Data type	Description
CollectionSpecimenID	int	Refers to ID of CollectionEvent (= Foreign key and part of primary key)
CollectorsName	nvarchar (255)	Name of the Collector
CollectorsAgentURI	varchar (255)	The URI of the Agent, e.g. as stored within the module DiversityAgents
CollectorsSequence	datetime	The order of collectors in a team. Automatically set by the database system *Default value: getdate()*
CollectorsNumber	nvarchar (50)	Number assigned to a specimen or a batch of specimens by the collector during the collection event (= 'field number')
Notes	nvarchar (MAX)	Notes about the collector, e.g. if the name is uncertain
DataWithholdingReason	nvarchar (255)	If the dataset is withhold, the reason for withholding the data, otherwise null
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this

		dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionEvent

The collection event where the specimen was collected

Column	Data type	Description
CollectionEventID	int	Unique ID for the collection event (= Primary key)
Version	int	The version of the dataset. Automatically set by the system. *Default value: (1)*
SeriesID	int	The ID of the related expedition. Relates to the PK of the table CollectionExpedition (Foreign key).
CollectorsEventNumber	nvarchar (50)	Number assigned to a collection event by the collector (= 'field number')
CollectionDate	datetime	The date of the event calulated from the entries in CollectionDay, -Month and -Year.
CollectionDay	tinyint	The day of the date of the event or when the collection event started
CollectionMonth	tinyint	The month of the date of the event or when the collection event started
CollectionYear	smallint	The year of the date of the event or when the collection event started
CollectionDateSupplement	nvarchar (100)	Verbal or additional collection date information, e.g. 'end of summer 1985', 'first quarter', '1888-1892'. The end date if the collection event comprises a period. The time of the event if necessary.
CollectionDateCategory	nvarchar (50)	Category of the date of the identification e.g. "system", "estimated" (= foreign key, see in table CollEventDateCategory_Enum)
CollectionTime	varchar (50)	The time of the event or when the collection event started

CollectionTimeSpan	varchar (50)	The time span e.g. in seconds of the collection event
LocalityDescription	nvarchar (MAX)	Locality description of the locality, exactly as written on the original label (i.e. without corrections during data entry)
HabitatDescription	nvarchar (MAX)	Geo-ecological description of the locality, exactly as written on the original label (i.e. without corrections during data entry)
ReferenceTitle	nvarchar (255)	The title of the publication where the collection event was published. Note this is only a cached value where ReferenceURI is present
ReferenceURI	varchar (255)	URI (e.g. LSID) of the source publication where the collection event is published, may e.g. refer to the module DiversityReferences
CollectingMethod	nvarchar (MAX)	Description of the method used for collecting the samples, e.g. traps, moist chambers, drag net
Notes	nvarchar (MAX)	Notes about the collection event
CountryCache	nvarchar (50)	The country where the collection event took place. Cached value derived from an entry in CollectionGeography
DataWithholdingReason	nvarchar (255)	If the dataset is withhold, the reason for withholding the data, otherwise null
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionEventImage

The images showing the site of the collection event

Column	Data type	Description
CollectionEventID	int	Unique ID for the collection event (= Primary key)

URI	varchar (255)	The complete URI address of the image. This is only a cached value if ResourceID is available referring to the module DiversityResources
ResourceURI	varchar (255)	The URI of the resource (e.g. see module DiversityResources)
ImageType	nvarchar (50)	Type of the image, e.g. map
Notes	nvarchar (MAX)	Notes to this image concerning the collection event
DataWithholdingReason	nvarchar (255)	If the dataset is withhold, the reason for withholding the data, otherwise null
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionEventLocalisation

The geographic localisation of a collection event

Column	Data type	Description
CollectionEventID	int	Refers to the ID of CollectionEvent (= Foreign key and part of primary key)
LocalisationSystemID	int	Refers to the ID of LocalisationSystem (= Foreign key and part of primary key)
Location1	nvarchar (255)	Either a named location selected from a thesaurus (e. g. 'Germany, Bavaria, Kleindingharting') or altitude range or other values (e. g. 100-200 m)
Location2	nvarchar (255)	Corresponding value to Location1 e.g. ID or URI of gazetteer or thesaurus
LocationAccuracy	nvarchar (50)	The accuracy of the determination of this locality
LocationNotes	nvarchar (MAX)	Notes on the location
DeterminationDate	smalldatetime	Date of the determination of the geographical localisation

DistanceToLocation	varchar (50)	Distance from the specified place to the real location of the collection event (m)
DirectionToLocation	varchar (50)	Direction from the specified place to the real location of the collection event (Degrees rel. to north)
ResponsibleName	nvarchar (255)	The name of the agent (person or organization) responsible for this entry. *Default value: [dbo].[CurrentUserName]()*
ResponsibleAgentURI	varchar (255)	URI of the person or organisation responsible for the data (see e.g. module DiversityAgents)
AverageAltitudeCache	real	Calculated altitude as parsed from the location fields
AverageLatitudeCache	real	Calculated latitude as parsed from the location fields
AverageLongitudeCache	real	Calculated longitude as parsed from the location fields
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionEventProperty

A property of a collection site, e.g. exposition, slope, vegetation. May refer to Diversity Workbench module DiversityScientificTerms

Column	Data type	Description
CollectionEventID	int	Refers to the ID of CollectionEvent (= Foreign key and part of primary key)
PropertyID	int	The ID of the descriptor of the collection event, foreign key, see table Descriptor
DisplayText	nvarchar (255)	The text for the property as shown e.g. in a user interface
PropertyURI	varchar (255)	URI referring to an external datasource e.g. DiversityTerminology
PropertyHierarchyCache	nvarchar (MAX)	A cached text of the

		complete name of the descriptor including superior categories if present
PropertyValue	nvarchar (255)	The value of a captured feature e.g. temperature, pH, vegetation etc. If there is a range this is the lower or first value
ResponsibleName	nvarchar (255)	The name of the agent (person or organization) responsible for this entry. *Default value: [dbo].[CurrentUserName]()*
ResponsibleAgentURI	varchar (255)	URI of the person or organisation responsible for the data (see e.g. module DiversityAgents)
Notes	nvarchar (MAX)	Notes about the property of the colletion site.
AverageValueCache	float	For numeric values - a cached average value according to the
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionEventSeries

The series whithin which collection events take place

Column	Data type	Description
SeriesID	int	Primary key. The ID for this expedition (= Primary key)
SeriesParentID	int	The ID of the superior expedition
Description	nvarchar (MAX)	The description of the expedition as it will be printed on e.g. the label
SeriesCode	nvarchar (50)	The user defined code for an expedition
Notes	nvarchar (MAX)	Notes about this expedition
DateCache	datetime	The first date of the depending events, used for sorting the expeditions

		[controlled by the database]
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionEventSeriesImage

The images showing the sites of a collection event series, e.g. an expedition or sampling plot

Column	Data type	Description
SeriesID	int	Unique ID for the collection event series (= Foreign key and part of primary key)
URI	varchar (255)	The complete URI address of the image. This is only a cached value if ResourceID is available referring to the module DiversityResources
ResourceURI	varchar (255)	The URI of the resource (e.g. see module DiversityResources)
ImageType	nvarchar (50)	Type of the image, e.g. map
Notes	nvarchar (MAX)	Notes to this image concerning the collection event
DataWithholdingReason	nvarchar (255)	If the dataset is withhold, the reason for withholding the data, otherwise null
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionExternalDatasource

CollectionExternalDatasource document the sources of the names.

Column	Data type	Description
ExternalDatasourceID	int	An ID to identify an external data collection of collection specimen (primary key, the ID has no meaning outside of the DiversityWorkbench system)
ExternalDatasourceName	nvarchar (255)	The name of the data collection that has been integrated or can be linked to for further analysis
ExternalDatasourceVersion	nvarchar (255)	The version of this data collection (either official version number, or dates when the collection was integrated)
Rights	nvarchar (500)	A description of copyright agreements or permission to use data from the external database
ExternalDatasourceAuthors	nvarchar (200)	The persons or institutions responsible for the external database
ExternalDatasourceURI	nvarchar (300)	The URI of the database provider resp. the external database
ExternalDatasourceInstitution	nvarchar (300)	The institution responsible for the external database
InternalNotes	nvarchar (1500)	Additional notes concerning this data collection
ExternalAttribute_NameID	nvarchar (255)	The table and field name in the external data collection to which CollectionExternalID refers
PreferredSequence	tinyint	For selection in e.g. picklists: of several equal names only the name from the source with the lowest preferred sequence will be provided.
Disabled	bit	If this source should be disabled for selection of names e.g. in picklists

Table CollectionManager

Collection managers within DiversityCollection, responsible of specimen transactions

Column	Data type	Description

LoginName	nvarchar (50)	A login name which the user uses for access the DivesityWorkbench, Microsoft domains, etc..
AdministratingCollectionID	int	ID for the collection for which the Manager has the right to administrate the transaction. Corresponds to AdministratingCollectionID in table Transaction.

Table CollectionProject

The projects within which the collection specimen were placed

Column	Data type	Description
CollectionSpecimenID	int	Refers to the ID of CollectionSpecimen (= Foreign key and part of primary key)
ProjectID	int	ID of the project to which the specimen belongs (Projects are defined in DiversityProjects)
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionRequester

Requesters within DiversityCollection, responsible of specimen transactions

Column	Data type	Description
LoginName	nvarchar (50)	A login name which the user uses for access to the DivesityWorkbench, Microsoft domains, etc..
AdministratingCollectionID	int	ID for the collection for which the Requester has the right to request specimen. Corresponds to AdministratingCollectionID in table Transaction.
IncludeSubcollections	bit	If the subcollections of the

		administrating collection are accessible for a request

Table CollectionSpecimen

The data directly attributed to the collection specimen

Column	Data type	Description
CollectionSpecimenID	int	Unique reference ID for the collection specimen record (primary key)
Version	int	The version of the dataset *Default value: (1)*
CollectionEventID	int	Refers to the ID of CollectionEvent (= Foreign key and part of primary key)
CollectionID	int	ID of the Collection as stored in table Collection (= foreign key, see table Collection)
AccessionNumber	nvarchar (50)	Accession number of the specimen within the collection, e.g. "M-29834752"
AccessionDate	datetime	The date of the accession calculated from the entries in AccessionDay, -Month and -Year
AccessionDay	tinyint	The day of the date when the specimen was acquired in the collection
AccessionMonth	tinyint	The month of the date when the specimen was acquired in the collection
AccessionYear	smallint	The year of the date when the specimen was acquired in the collection
AccessionDateSupplement	nvarchar (255)	Verbal or additional accession date information, e.g. 'end of summer 1985', 'first quarter', '1888-1892'
AccessionDateCategory	nvarchar (50)	Category of the date of the identification e.g. "system", "estimated" (= foreign key, see in table xColl_DateCategory_Enum)
DepositorsName	nvarchar (255)	The name of the depositor(s) (person or organization responsible for deposition). Where entire collections are deposited, this should also contain the collection name (e.g. 'Herbarium P. Döbbler')
DepositorsAgentURI	varchar (255)	The URI of the depositor(s) (person or organization

		responsible for deposition)
DepositorsAccessionNumber	nvarchar (50)	Accession number of the specimen within the previous or original collection, e.g. 'D-23948'
LabelTitle	nvarchar (255)	The title of the label e.g. for printing labels.
LabelType	nvarchar (50)	Printed, typewritten, typewritten with handwriting added, entirely in handwriting, etc.
LabelTranscriptionState	nvarchar (50)	The state of the transcription of a label into the database: 'Not started', 'incomplete', 'complete'
LabelTranscriptionNotes	nvarchar (255)	User defined notes concerning the transcription of the label into the database
ExsiccataURI	varchar (255)	If specimen is an exsiccata: The URI of the Exsiccata series, e.g. as stored within the DiversityExsiccata module
ExsiccataAbbreviation	nvarchar (255)	If specimen is an exsiccata: Standard abbreviation of the exsiccata (not necessarily a unique identifier; editors or publication places may change over time)
OriginalNotes	nvarchar (MAX)	Notes found on the label of the specimen, by the original collector or from a later revision
AdditionalNotes	nvarchar (MAX)	Additional notes made by the editor of the specimen record, e. g. 'doubtful identification/locality'
ReferenceTitle	nvarchar (255)	The title of the publication where the specimen was published. Note this is only a cached value where ReferenceURI is present
ReferenceURI	varchar (255)	URI (e.g. LSID) of reference where specimen is published, e.g. referring to the module DiversityReferences
Problems	nvarchar (255)	Description of a problem that occurred during data editing. Typically these entries should be deleted after help has been obtained. Do not enter scientific problems here; use AdditionalNotes for such permanent problems!
DataWithholdingReason	nvarchar (255)	If the dataset is withhold, the reason for withholding the

		data, otherwise null
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*
InternalNotes	nvarchar (MAX)	Internal notes that should not be published e.g. on websites
ExternalDatasourceID	int	An ID to identify an external data collection of collection specimen (primary key, the ID has no meaning outside of the DiversityWorkbench system)
ExternalIdentifier	nvarchar (100)	The identifier of the external specimen as defined in the external datasource

Table CollectionSpecimenImage

The images of a collection specimen or of an identification unit within this specimen

Column	Data type	Description
CollectionSpecimenID	int	Refers to the ID of CollectionSpecimen (= Foreign key and part of primary key)
URI	varchar (255)	The complete URI address of the image. This is only a cached value if ResourceID is available referring to the module DiversityResources
ResourceURI	varchar (255)	The URI of the image, e.g. as stored in the module DiversityResources.
SpecimenPartID	int	Optional: If the dataset is not related to a part of a specimen, the ID of a related part (= foreign key)
IdentificationUnitID	int	If image refers to only on out of several identification units for a specimen, refers to the ID of an IdentificationUnit for a CollectionSpecimen (= foreign key)
ImageType	nvarchar (50)	Type of the image, e.g. label
Notes	nvarchar (MAX)	Notes about the specimen

		image
DataWithholdingReason	nvarchar (255)	If the dataset is withhold, the reason for withholding the data, otherwise null
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionSpecimenPart

Parts of a collection specimen. Includes a possible hierarchy of the parts

Column	Data type	Description
CollectionSpecimenID	int	Refers to the ID of CollectionSpecimen (= Foreign key and part of primary key)
SpecimenPartID	int	ID for a part of a specimen (part of primary key) *Default value: (1)*
DerivedFromSpecimenPartID	int	SpecimenPartID of the specimen from which the current specimen is derived from
PreparationMethod	nvarchar (MAX)	The method used for the preparation of the part of the specimen, e.g. the inoculation method for cultures
PreparationDate	datetime	The date and time when the part was preparated e.g when it was separated from the source object
AccessionNumber	nvarchar (50)	Accession number of the part of the specimen within the collection if it is different from the accession number of the specimen as stored in the table CollectionSpecimen, e.g. "M-29834752"
PartSublabel	nvarchar (50)	The label for a part of a specimen, e.g. if duplicats of a specimen have a separate number
CollectionID	int	ID of the Collection as stored in table Collection (= foreign key, see table Collection)

MaterialCategory	nvarchar (50)	Material category of specimen. Examples: 'herbarium sheets', 'drawings', 'microscopic slides' etc. (= foreign key, see table CollMaterialCategory_Enum) *Default value: N'specimen'*
StorageLocation	nvarchar (255)	A code identifying the place where the specimen is stored within the collection. Frequently the accepted scientific name is used as storage location code.
Stock	tinyint	Number of stock units if the specimen is stored in separated units e.g. several boxes or vessels (max. 255)
Notes	nvarchar (MAX)	Notes concerning the storage of the sample
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionSpecimenProcessing

The processing that was applied to a collection specimen

Column	Data type	Description
CollectionSpecimenID	int	Refers to ID of CollectionSpecimen (= Foreign key and part of primary key)
ProcessingDate	datetime	Date and time of the start of the processing *Default value: getdate()*
ProcessingID	int	ID of the processing. Refers to ProcessingID in table Processing (foreign key) *Default value: (1)*
Protocoll	nvarchar (100)	The label of the processing protocoll
SpecimenPartID	int	Optional: If the dataset is related to a part of a specimen, the ID of a related part (= foreign key, see table CollectionSpecimenPart)

ProcessingDuration	varchar (50)	The duration of the processing including the unit (e.g. 5 min) or the end of the processing starting at the processingDate (e.g. 23.05.2008)
ResponsibleName	nvarchar (255)	Name of the person or institution responsible for the determination *Default value: [dbo].[CurrentUserName]()*
ResponsibleAgentURI	varchar (255)	URI of the person or institution responsible for the determination (= foreign key) as stored in the module DiversityAgents.
Notes	nvarchar (MAX)	Notes about the processing
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionSpecimenRelation

The relations of a collection specimen to other collection specimen

Column	Data type	Description
CollectionSpecimenID	int	Unique reference ID for the collection specimen record (primary key)
RelatedSpecimenURI	varchar (255)	URI of the related specimen
RelatedSpecimenDisplayText	varchar (255)	The name of a related specimen as shown e.g. in a user interface
RelationType	nvarchar (50)	Type of the relation between the specimen (= foreign key, see table CollRelationType_Enum)
RelatedSpecimenCollectionID	int	ID of the Collection as stored in table Collection (= foreign key, see table Collection)
RelatedSpecimenDescription	nvarchar (MAX)	Description of the related specimen
Notes	nvarchar (MAX)	Notes on the relation to the specimen

IsInternalRelationCache	bit	If the relation represents a connection between specimen in this database *Default value: (1)*
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table CollectionSpecimenTransaction

The transactions in which a specimen was involved

Column	Data type	Description
CollectionSpecimenID	int	Refers to ID of CollectionSpecimen (= Foreign key and part of primary key)
TransactionID	int	Unique ID for the transaction (= Foreign key and part of primary key)
SpecimenPartID	int	Optional: If the dataset is related to a part of a specimen, the ID of a related part (= foreign key, see table CollectionSpecimenPart)
IsOnLoan	bit	True if a specimen is on loan
LogInsertedBy	nvarchar (50)	Name of user who first entered (typed or imported) the data. *Default value: user_name()*
LogInsertedWhen	smalldatetime	Date and time when the data were first entered (typed or imported) into this database. *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Name of user who last updated the data. *Default value: user_name()*
LogUpdatedWhen	smalldatetime	Date and time when the data were last updated. *Default value: getdate()*

Table ExternalRequestCredentials

External requestors with the permission to create a request for a loan

Column	Data type	Description
RequesterLogin	nvarchar (50)	Login of the person responsible for the loan requests in the collection
AdministratingCollectionID	int	The ID of the collection which gets the request for a loan. Corresponds to the AdministratingCollectionID in table Transaction.
RequestingCollectionID	int	The ID of the collection for which the requester has the permission to create a request

Table Identification

The identifications of the organisms within a specimen

Column	Data type	Description
CollectionSpecimenID	int	Refers to the ID of CollectionSpecimen (= Foreign key and part of primary key)
IdentificationUnitID	int	Refers to the ID of IdentficationUnit (= foreign key and part of primary key)
IdentificationSequence	smallint	The sequence of the identifications. The last identification (having the highest sequence) is regarded as valid *Default value: (1)*
IdentificationDate	datetime	The date of the identification calculated from the entries in IdentificationDay, -Month and -Year
IdentificationDay	tinyint	The day of the identification event
IdentificationMonth	tinyint	The month of the identification event
IdentificationYear	smallint	The year of the identification event. The year may be empty if only the day or month are known.
IdentificationDateSupplement	nvarchar (255)	Verbal or additional identification date information, e.g. 'end of summer 1985', 'first quarter', '1888-1892'
IdentificationDateCategory	nvarchar (50)	Category of the date of the identification e.g. "system", "estimated" (= foreign key, see in table CollDateCategory_Enum)

VernacularTerm	nvarchar (255)	Name or term other than a taxonomic (= scientific) name, e.g. 'pine', 'limestone', 'conifer', 'hardwood'
TaxonomicName	nvarchar (255)	Valid name of the species (including the taxonomic author where available. Example: 'Rosa canina L.'
NameURI	varchar (255)	The URI of the taxonomic name, e.g. as provided by the module DiversityTaxonNames.
IdentificationCategory	nvarchar (50)	Category of the identification e.g. 'determination', 'confirmation', 'absence' (= foreign key, see table CollIdentificationCategory_Enum)
IdentificationQualifier	nvarchar (50)	Qualification of the identification e.g. "cf."," aff.", "sp. nov." (= foreign key, see table CollIdentificationQualifier_Enum)
TypeStatus	nvarchar (50)	If identification unit is type of a taxonomic name: holotype, syntype, etc. (= foreign key, see table CollTypeStatus_Enum)
TypeNotes	nvarchar (MAX)	Notes concerning the typification of this specimen
ReferenceTitle	nvarchar (255)	Publications or authoritative opinions of scientist used during the identification process. Example: enter 'Schmeil-Fitschen 1995' if this field flora was used.
ReferenceURI	varchar (255)	The URI of the reference e.g. as provided by the module DiversityReferences
Notes	nvarchar (MAX)	User defined notes, e.g. the reason for a re-determination / change of the name, etc.
ResponsibleName	nvarchar (255)	Name of the person or institution responsible for the determination *Default value: [dbo].[CurrentUserName]()*
ResponsibleAgentURI	varchar (255)	URI of the person or institution responsible for the determination (= foreign key) as stored in the module DiversityAgents.
LogCreatedWhen	datetime	Date and time when the dataset was created *Default value: getdate()*

LogCreatedBy	nvarchar (50)	Login of the user who created the dataset *Default value: user_name()*
LogUpdatedWhen	datetime	Date and time when the dataset was changed *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Login of the user who changed the dataset *Default value: user_name()*

Table IdentificationUnit

Organism that is present in or on a collection specimen

Column	Data type	Description
CollectionSpecimenID	int	Refers to the ID of CollectionSpecimen (= Foreign key and part of primary key)
IdentificationUnitID	int	ID of the identification unit (= part of Primary key). Usually one of possibly several organisms present on the collection specimen. Example: parasite with hyperparasite on plant leaf = 3 units,
LastIdentificationCache	nvarchar (255)	The last identification as entered in table Identification
FamilyCache	nvarchar (255)	A cached value of the family of the taxon of the last identification. Can be set by the editor if NameURI in table Identification is NULL, otherwise set by the system.
OrderCache	nvarchar (255)	A cached value of the order of the taxon of the last identification. Can be set by the editor if NameURI in table Identification is NULL, otherwise set by the system.
TaxonomicGroup	nvarchar (50)	Taxonomic group the organism identified by this unit belongs to. Groups listed in table CollTaxonomicGroup_Enum (= foreign key)
OnlyObserved	bit	True if the organism was only observed rather than collected. It is therefore not present on the preserved specimen. Example: Tree under which the collected mycorrhizal fungus grew. *Default value: (0)*
RelatedUnitID	int	The IdentificationUnitID of the organism or substrate, on

		which this organism is growing (= foreign key)
RelationType	nvarchar (50)	The relation of an unit to its substrate, e.g. parasitism, symbiosis etc. as stored in CollRelationType_Enum (= foreign key)
ColonisedSubstratePart	nvarchar (255)	If a substrate association exists: part of the substrate that is affected in the interaction (e.g. 'leaves' if a fungus is growing on the leaves of an infected plant)
LifeStage	nvarchar (255)	Examples: 'II, III' for spore generations of rusts or 'seed', 'seedling' etc. for higher plants
Gender	nvarchar (50)	The gender of the identification unit, e.g. 'male'
NumberOfUnits	smallint	The number of units of this identification unit, e.g. 400 beetle in a bottle
ExsiccataNumber	nvarchar (50)	If specimen is an exsiccata: Number of current specimen within the exsiccata series
ExsiccataIdentification	smallint	Refers to the IdentificationSequence in Identification (= foreign key). The name under which the collection specimen resp. this unit is published within an exsiccata.
UnitIdentifier	nvarchar (50)	An identifier for the identification of the unit e.g. a number painted on a tree within an experimental plot
UnitDescription	nvarchar (50)	Description of the unit, esp. if not the an organism but parts or remnants of it were present or observed, e.g. a nest of an insect or a song of a bird
Circumstances	nvarchar (50)	Circumstances of the occurence of the organism
DisplayOrder	smallint	The sequence in which the units within this specimen will appear on e.g. a label where the first unit may be printed in the header and others in the text below. 0 means the unit should not appear on a label. *Default value: (1)*
Notes	nvarchar (MAX)	Further information on the identification unit or interaction, e. g. infection symptoms like 'producing galls'

LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table IdentificationUnitAnalysis

The analysis values taken from an identification unit

Column	Data type	Description
CollectionSpecimenID	int	Refers to the ID of CollectionSpecimen (= Foreign key and part of primary key)
IdentificationUnitID	int	Refers to the ID of IdentficationUnit (= foreign key and part of primary key)
AnalysisID	int	Analysis ID, foreign key of table Analysis.
AnalysisNumber	nvarchar (50)	Number of the analysis
AnalysisResult	nvarchar (MAX)	The result of the analysis
ExternalAnalysisURI	varchar (255)	An URI for an analysis as defined in an external datasoure
ResponsibleName	nvarchar (255)	Name of the person or institution responsible for the determination *Default value: [dbo].[CurrentUserName]()*
ResponsibleAgentURI	varchar (255)	URI of the person or institution responsible for the determination (= foreign key) as stored in the module DiversityAgents.
AnalysisDate	nvarchar (50)	The date of the analysis
SpecimenPartID	int	ID of the part of a specimen (optional, Foreign key) if the analysis was done with a part of the specimen (see table CollectionSpecimenPart).
Notes	nvarchar (MAX)	Notes concerning this analysis
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*

LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table IdentificationUnitInPart

The list of the organisms that are found in a part of the specimen

Column	Data type	Description
CollectionSpecimenID	int	Refers to the ID of CollectionSpecimen (= Foreign key and part of primary key)
IdentificationUnitID	int	ID of the identification unit (= part of Primary key). Usually one of possibly several organisms present on the collection specimen. Example: parasite with hyperparasite on plant leaf = 3 units,
SpecimenPartID	int	ID of the part of a specimen (optional, Foreign key) if the identification unit is located on a part of the specimen (see table CollectionSpecimenPart).
DisplayOrder	smallint	The sequence in which the units within this part will appear on e.g. a label where the first unit may be printed in the header and others in the text below. 0 means the unit should not appear on a label. *Default value: (1)*
Description	nvarchar (500)	A description of the unit, esp. if not a whole unit but e.g. parts of it are stored in the collection, e.g. a nest of a bird
LogInsertedBy	nvarchar (50)	Name of user who first entered (typed or imported) the data. *Default value: user_name()*
LogInsertedWhen	smalldatetime	Date and time when the data were first entered (typed or imported) into this database. *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Name of user who last updated the data. *Default value: user_name()*
LogUpdatedWhen	smalldatetime	Date and time when the data were last updated.

		Default value: getdate()

Table LocalisationSystem

The geographic localisation systems, e.g. coordinates

Column	Data type	Description
LocalisationSystemID	int	Unique ID for the localisation system (= Primary key)
LocalisationSystemParentID	int	LocalisationSystemID of the superior LocalisationSystem
LocalisationSystemName	nvarchar (100)	Name of the system used for the determination of the place of the collection, e. g. Gauss-Krüger, MTB, GIS
DefaultAccuracyOfLocalisation	nvarchar (50)	The default for the accuracy of values that can be reached with this method
DefaultMeasurementUnit	nvarchar (50)	The default measurement unit for the localisation system, e.g. m, geograpic coordinates
ParsingMethodName	nvarchar (50)	Internal value, specifying a programming method used for parsing text in fields Location1/Location2 in table CollectionLocalisation
DisplayText	nvarchar (50)	Short abbreviated description of the localisation system as displayed in the user interface
DisplayEnable	bit	Specifies if this item is enabled to be used within the database. LocalisationSystems can be disabled to avoid seeing them, but to keep the definition for the future.
DisplayOrder	smallint	The order in which the entries are displayed. The order may be changed at any time, but all values must be unique.
Description	nvarchar (255)	Description of the localisation method
DisplayTextLocation1	nvarchar (50)	Short abbreviated description of the attribute Location1 in the table CollectionGeography as displayed in the user interface
DescriptionLocation1	nvarchar (255)	Description of the attribute Location1 in the table CollectionGeography as displayed in the user interface
DisplayTextLocation2	nvarchar (50)	Short abbreviated description of the attribute Location2 in

		the table CollectionGeography as displayed in the user interface
DescriptionLocation2	nvarchar (255)	Description of the attribute Location2 in the table CollectionGeography as displayed in the user interface

Table Processing

The processings of the specimen

Column	Data type	Description
ProcessingID	int	ID of the processing (Primary key)
ProcessingParentID	int	The ID of the superior type of the processing
DisplayText	nvarchar (50)	The display text of the processing as shown e.g. in a user interface
Description	nvarchar (MAX)	Description of the processing
Notes	nvarchar (MAX)	Notes about the processing
ProcessingURI	varchar (255)	An URI for a processing as defined in an external datasource
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table ProcessingMaterialCategory

The processings that are possible for a certain material category

Column	Data type	Description
ProcessingID	int	ID of the processing. Refers to ProcessingID in table Processing (foreign key) *Default value: (1)*
MaterialCategory	nvarchar (50)	Material category of specimen. Examples: 'herbarium sheets', 'drawings', 'microscopic slides' etc.

		Default value: N'specimen'
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table Property

The list of the properties that can be specified for the collection site

Column	Data type	Description
PropertyID	int	Unique ID for the localisation system (= Primary key)
PropertyParentID	int	LocalisationSystemID of the superior LocalisationSystem
PropertyName	nvarchar (100)	Name of the system used for the determination of the place of the collection, e. g. Gauss-Krüger, MTB, GIS
DefaultAccuracyOfProperty	nvarchar (50)	The default for the accuracy of values that can be reached with this method
DefaultMeasurementUnit	nvarchar (50)	-
ParsingMethodName	nvarchar (50)	Internal value, specifying a programming method used for parsing text in fields Location1/Location2 in table CollectionLocalisation
DisplayText	nvarchar (50)	Short abbreviated description of the localisation system as displayed in the user interface
DisplayEnabled	bit	Specifies if this item is enabled to be used within the database. LocalisationSystems can be disabled to avoid seeing them, but to keep the definition for the future.
DisplayOrder	smallint	The order in which the entries are displayed. The order may be changed at any time, but all values must be unique.
Description	nvarchar (255)	Description of the localisation method

Table Transaction

Transactions like loan, borrow, gift, exchange etc. of specimen if they are

e.g. permanently or temporary transfered from one collection to another

Column	Data type	Description
TransactionID	int	Unique ID for the transaction (= Primary key)
ParentTransactionID	int	The ID of a preceeding transaction of a superior transaction if transactions are organized in a hierarchy
TransactionType	nvarchar (50)	Type of the transaction e.g. gift in or out, exchange in or out, purchase in or out *Default value: N'exchange'*
TransactionTitle	nvarchar (200)	The title of the transaction as e.g. shown in an user interface
ReportingCategory	nvarchar (50)	A group defined for the transaction, e. g. a taxonomic group as used for exchange balancing
AdministratingCollectionID	int	ID of the collection thas is responsible for the administration of the transaction.
MaterialDescription	nvarchar (MAX)	ID of the project to which the transaction belongs (Projects are defined in DiversityProjects) *Default value: ''*
MaterialCategory	nvarchar (50)	Material category of specimen. Examples: 'herbarium sheets', 'drawings', 'microscopic slides' etc. *Default value: N'specimen'*
MaterialCollectors	nvarchar (MAX)	The collectors of the material
FromCollectionID	int	The ID of the collection from which the specimen were transfered, e.g. the donating collection of a gift
FromTransactionPartnerName	nvarchar (255)	Name of the person or institution from which the specimen were transfered, e.g. the donator of a gift
FromTransactionPartnerAgentURI	varchar (255)	The URI of the transaction partner (see e.g. module DiversityAgents)
FromTransactionNumber	nvarchar (50)	Number or code by which a transaction may be recorded by the administration of the source of the specimen, e.g. the donating collection of a gift
ToCollectionID	int	The ID of the collection to which the specimen were

		transfered, e.g. the receiver of a gift
ToTransactionPartnerName	nvarchar (255)	Name of the person or institution to which the specimen were transfered, e.g. the receiver of a gift
ToTransactionPartnerAgentURI	varchar (255)	The URI of the transaction partner (see e.g. module DiversityAgents)
ToTransactionNumber	nvarchar (50)	Number or code by which a transaction may be recorded by the administration of the destination of the specimen, e.g. the receiving collection of a gift
NumberOfUnits	smallint	The number of units that were (initially) included in the transaction
Investigator	nvarchar (50)	The investigator for whose study a transacted material was sent
TransactionComment	nvarchar (MAX)	Comments about the exchanged material addressed to the transaction partner
BeginDate	datetime	Date when the transaction started
AgreedEndDate	datetime	End of the transaction period, e.g. if the time for borrowing the specimen is restricted
ActualEndDate	datetime	Actual end of the transaction when e.g. the borrowed specimen were returned to the owner
InternalNotes	nvarchar (MAX)	Internal notes about this transaction, not to be published e.g. on a web page
ResponsibleName	nvarchar (255)	The person responsible for this transaction
ResponsibleAgentURI	varchar (255)	The URI of the person, team or organisation responsible for the data (see e.g. module DiversityAgents)
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

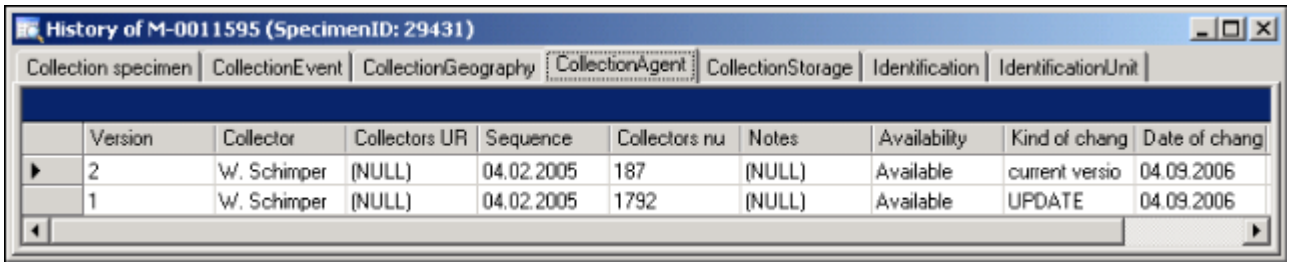
Table TransactionDocument

The history of transactions resp. the documents connected to the transactions

Column	Data type	Description
TransactionID	int	Unique ID for the Transaction, refers to table Transaction (= Part of primary key and foreign key)
Date	datetime	The date of the event of a transaction
TransactionText	nvarchar (MAX)	The text of a transaction document
TransactionDocument	image (2147483647)	A scanned document connected to this transaction event
InternalNotes	nvarchar (MAX)	Internal notes about this transaction event
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

History

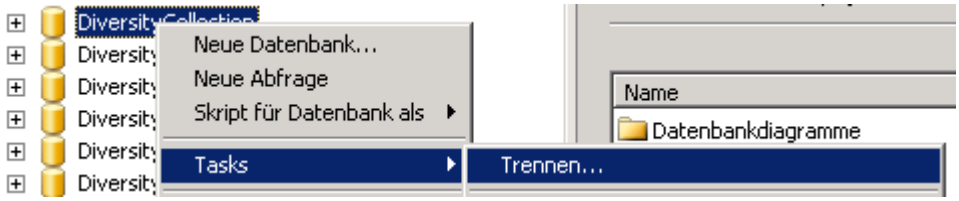
To inspect the history of a dataset click on the button. A window will open, showing all former states of the data in the tables with the current dataset at the top. The database DiversityCollection handles 2 different histories - one for the collection specimen and one for the collection event. The version shown in the header of the main window refers to these histories. The first number refers to the version of the specimen. If an event was defined a second number is shown, referring to the version of the collection event, e.g. 2 / 1 means version 2 of the specimen and version 1 of the collection event.



The version will be set automatically. If a dataset is changed the version will be increased if the last changes were done by a different user or the last change is more than 24 hours ago (for further details see topic Logging).

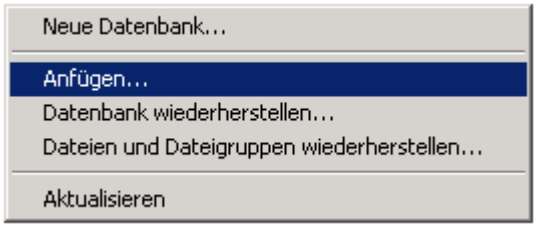
Backup

If you need to backup your database, you have to use the functionality provided by SQL-Server. To do this, you need administration rights in the database you want to create a backup. Open the Enterprise Manager for SQL-Server, choose the database and detach it from the server as shown in the image below.



After detaching the database, you can store a copy of the ..._Data.MDF File to keep it as a backup.

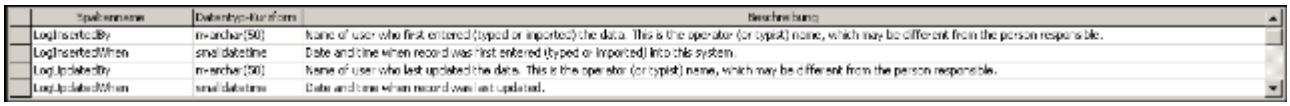
After storing the backup you have to attach the database.



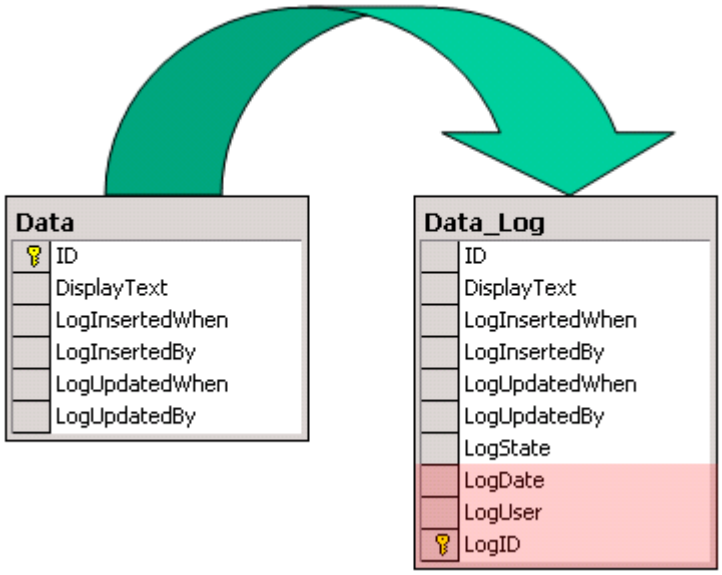
A dialog will appear where you have to select the original database file in your directory.

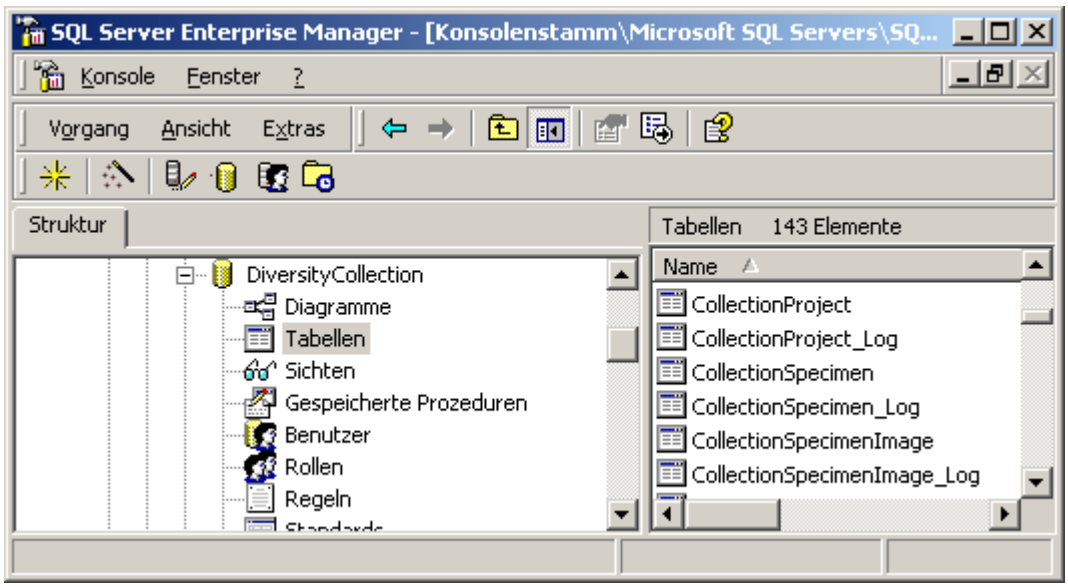
Logging

Changes within the database will be documented for each dataset together with the time and the responsible user in the columns shown in the image below.



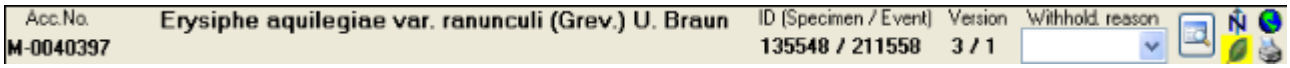
All main tables have a corresponding logging table. If you change or delete a dataset the orignial dataset will be stored in this logging table together with informations about who has done the changes and when it happend.





Version of datasets

The **Version** of a dataset consists of two parts, e.g. 3 / 1 as shown in the header of the specimen in the main form. The first part of the number refers to the version of the dataset as stored in table CollectionSpecimen. The second part refers to the version as stored in the table CollectionEvent. Both versions will be set to a higher number if the data in the tables themselves or in dependent tables are changed, e.g. the insertion of a new identification in the table Identification will increase the specimen part of the version from 3 to 4 resulting in a version 4 / 1 for the whole dataset. The changes in the version will only occur if the last changes in the data were more than **24 hours** ago or a **different user** is changing the data. This ensures, that a user can change several parts in a dataset within 24 hours and the version will only be increased by 1.



The data are stored in the tables CollectionSpecimen and CollectionEvent.

Database - Application tables

Informations concerning the application are stored in the application tables. The table ApplicationSearchSelectionStrings hold the SQL-strings of the predefined queries for the users. The table ApplicationEntityDescription holds the description of the database entities, if these are different from the description in the database, especially if other languages then English are to be supported.

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- ApplicationEntityDescription
- ApplicationSearchSelectionStrings
- ProjectProxy
- ProjectUser
- UserProxy

Table ApplicationEntityDescription

The description of the columns in different languages

Column	Data type	Description
TableName	varchar (50)	The name of the table within the database
ColumnName	varchar (50)	The name of the column of the table within the database
DisplayText	nvarchar (50)	The text for the column as shown e.g. in a user interface
Description	nvarchar (MAX)	The description of the content of the column
LanguageCode	nvarchar (50)	ISO 639: 2-letter codes for the language of DisplayText and Description
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table ApplicationSearchSelectionStrings

Selection strings for customized queries for users

Column	Data type	Description
UserName	varchar (50)	The name of the user who created this SQL string

		Default value: user_name()
SQLStringIdentifier	varchar (50)	The identifier for the selection string as shown in user interface
ItemTable	varchar (50)	The main table from which the datasets should be selected *Default value: 'TaxonName'*
SQLString	varchar (MAX)	SQL string for selecting datasets from the database
Description	nvarchar (MAX)	Description of the resultset and the purpose of the query
LogCreatedWhen	datetime	The time when this dataset was created *Default value: getdate()*
LogCreatedBy	nvarchar (50)	Who created this dataset *Default value: user_name()*
LogUpdatedWhen	datetime	The last time when this dataset was updated *Default value: getdate()*
LogUpdatedBy	nvarchar (50)	Who was the last to update this dataset *Default value: user_name()*

Table ProjectProxy

The projects as stored in the module DiversityProjects

Column	Data type	Description
ProjectID	int	ID of the project to which the specimen belongs (Projects are defined in DiversityProjects)
Project	nvarchar (50)	The name or title of the project as shown in a user interface (Projects are defined in DiversityProjects)

Table ProjectUser

The projects that a user can access

Column	Data type	Description
LoginName	nvarchar (50)	A login name which the user uses for access the DivesityWorkbench, Microsoft domains, etc..
ProjectID	int	ID of the project to which the specimen belongs (Projects are defined in DiversityProjects)

Table UserProxy

The user as stored in the module DiversityUsers

Column	Data type	Description
LoginName	nvarchar (50)	A login name which the user uses for access the DivesityWorkbench, Microsoft domains, etc..
CombinedNameCache	nvarchar (255)	The short name of the user,

		e.g. P. Smith *Default value: NULL*
UserURI	varchar (255)	URI of a user in a remote module, e.g. refering to UserInfo.UserID in database DiversityUsers

Diversity Workbench

The Diversity Workbench is composed of components for building and managing biodiversity information, each of which focuses on a particular domain. Each component can provide services to the other components. DiversityCollection can link data to the modules DiversityProjects, DiversityResources, DiversityExsiccatae, DiversityGazetteer, DiversityScientificTerms, DiversityTaxonNames, DiversityAgents, DiversityUsers and DiversityReferences as illustrated in the image below.

The modules communicate with each other to provide their services for the other modules.

Modules

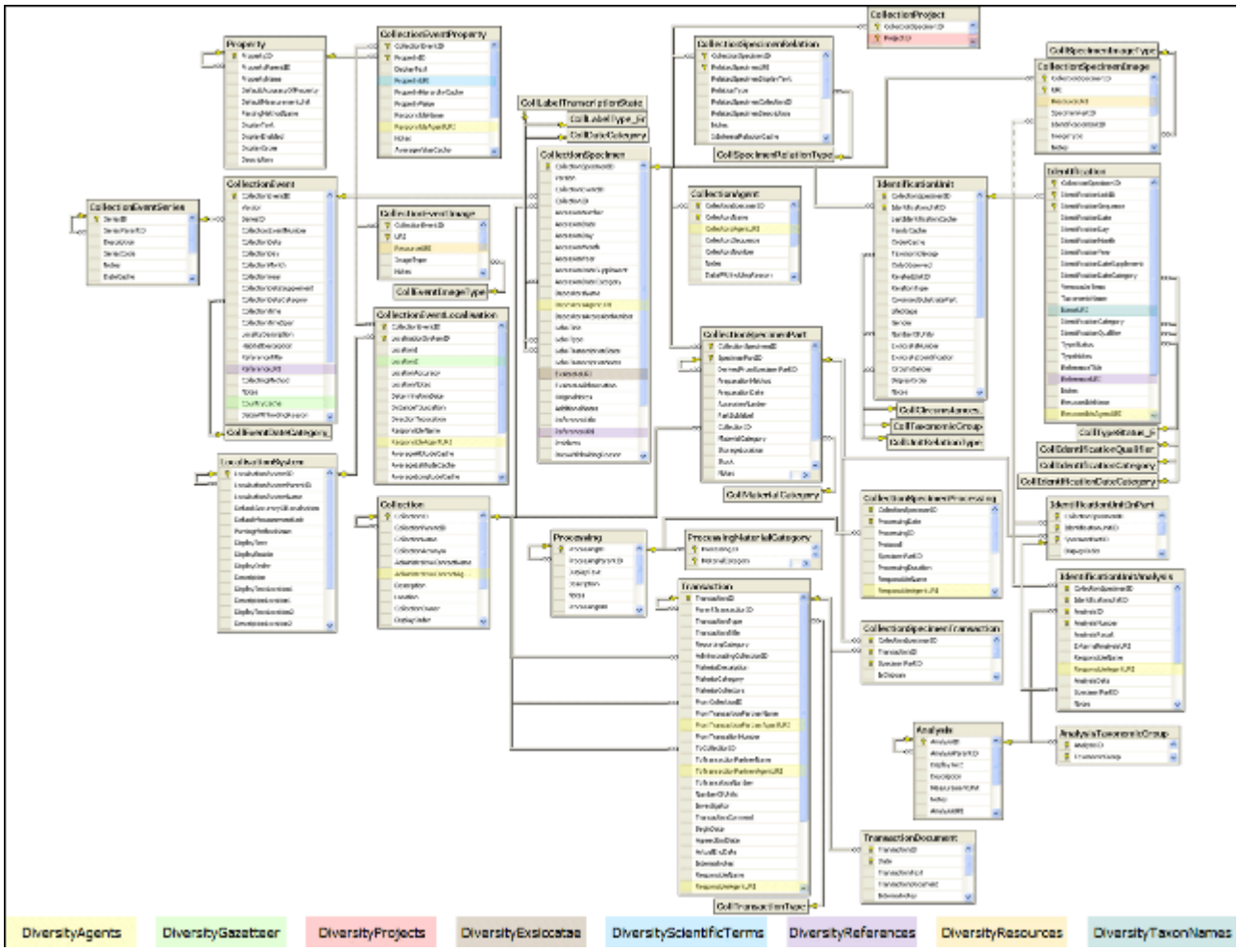
The Diversity Workbench is a set of components for building and managing biodiversity information, each of which focuses on a particular domain.

Diversity Agents	Administration of the agents, i.e. people and institutions which should be documented with e.g. their addresses
Diversity Collection	Administration of the scientific collections and specimens within these collections
Diversity Descriptions	Administration of descriptive data
Diversity Exsiccatae	Administration of exsiccatal series
Diversity Gazetteer	A data collection to enable the linking of geographical records with the Getty Thesaurus of Geographical Names (TGN), the Germ geographical names.
Diversity ScientificTerms	Data collections of scientific terms from foreign sources like vegetation, stratigraphy, soil science etc.
Diversity Projects	Administration of projects within the Diversity Workbench
Diversity References	Administration of references
Diversity Resources	Administration of resources like images, etc.
Diversity TaxonNames	Administration of taxonomic names, their synonyms and hierarchical position
Diversity Users	Administration of the users and their permissions within the Diversity Workbench

Each module provides services for the other modules. To use the service of a module, you need access to the database of the module and optionally the module application placed in

your application directory.

In the graphic below the connections to other modules within DiversityCollection are indicated.



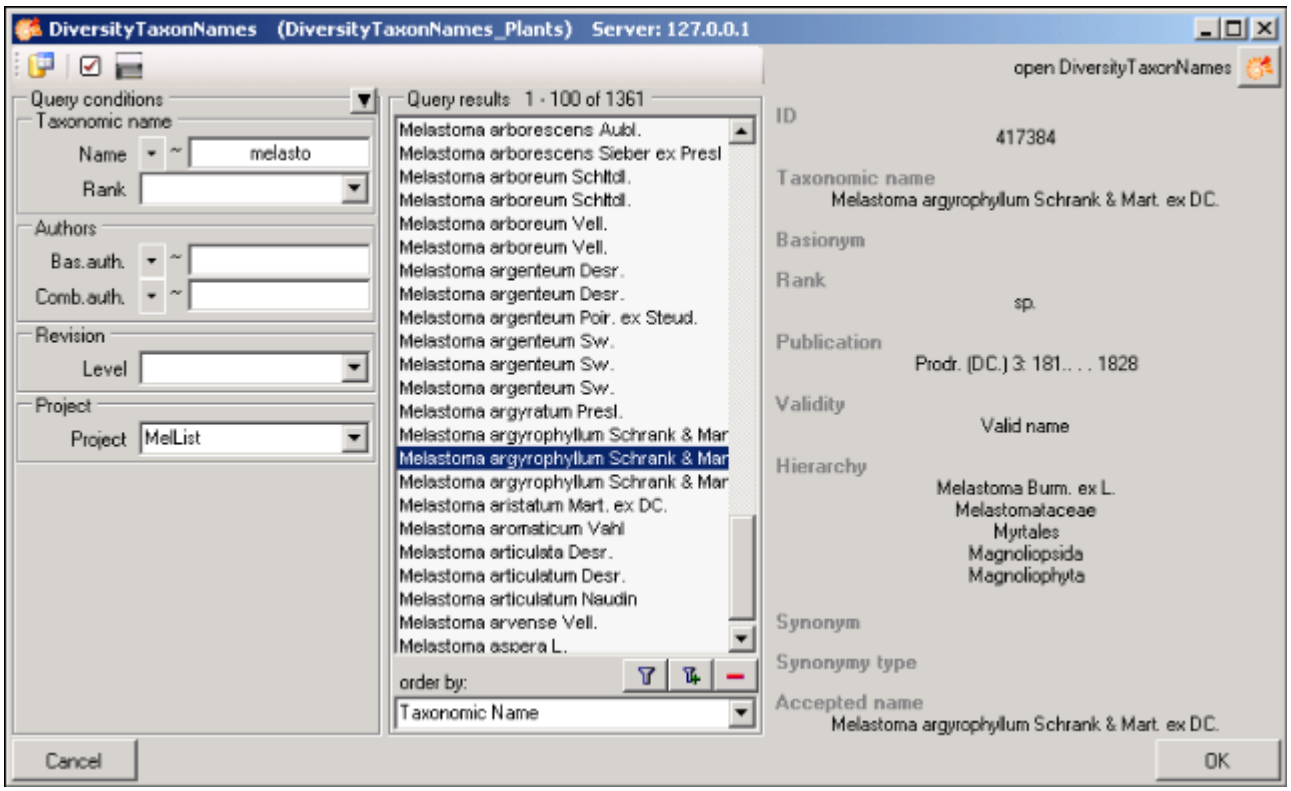
In the form a connection to a module of Diversity Workbench is a set of components for building and managing biodiversity information, each of which focuses on a particular domain.

Module related entry

The Diversity Workbench is a set of components for building and managing biodiversity information, each of which focuses on a particular domain. Each module provides services for the other modules. To use the service of a module, you need access to the database of the module and optionally the module application placed in your application directory. Entries related to an external module have a standard interface in the main form. There are 2 states of this interface:

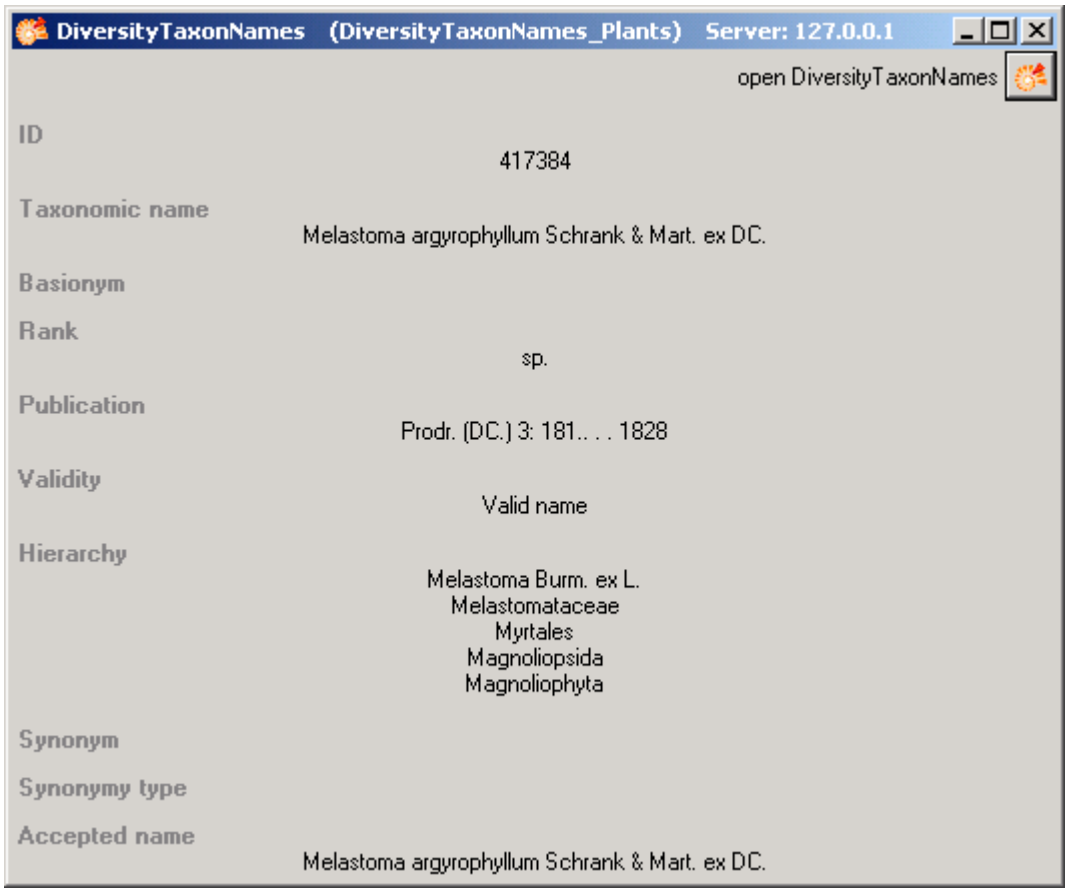
1 - the value is only set in the local database with no connection to the remote module

In this state, you can either type the value or select it from the values that are already available in the database. To get a list of the available values type the beginning of the value (you may use wildcards) and click on the button. If you want to set a relation to the remote module, click on the button. A window will open where you may select an entry from the foreign database.



2 - the value is related to the remote module

If the value has a relation to the remote module, the interface will appear as shown above. To release the connection to the remote module click on the button. If you need further information about the value, click on the button. This will open a form, showing an overview of the related value.



If the client application of the module is available you can inspect the details of the entry. To start the client application of the remote module, just click on the button.

3 - relation to a webservice

Some modules provide the possibility to link your data to an external webservice. For example DiversityTaxonNames gives you access to the taxonomic names of IndexFungorum. To establish a connection to an external webservice, click on the button. As with the link to modules within the Diversity Workbench a window will open where you can choose from either Diversity Workbench modules or external Webservices. See Webservice for further details

Resources

Images for specimen and the collection event can either be stored in DiversityCollection with their file path or in more detail in the module DiversityResources. For directly changing to DiversityResources click on the button.

For direct access to the resources in this module, you need the application **DiversityResources.exe** in your application directory, the database DiversityResources and a valid account in the database DiversityUsers. For more information see the Diversity Workbench Portal.

Reference

Details about References are stored in the module DiversityReferences. You can choose one of the entries in this module from the picklist. To directly change to DiversityReferences click on .

For access to the references from other modules, you need the application **DiversityReferences.exe** in your application directory. To use the application DiversityReferences.exe you need access to the database DiversityReferences. For more information see the Diversity Workbench Portal.

Webservice - foreign sources

Some modules within the Diversity Workbench provide the possibility to link your data to an external webservice. For example DiversityTaxonNames gives you access to the taxonomic names of IndexFungorum. To establish a connection to an external webservice, click on the button. A window will open where you can choose from either Diversity Workbench modules or external Webservices. The currently provided webservices are:

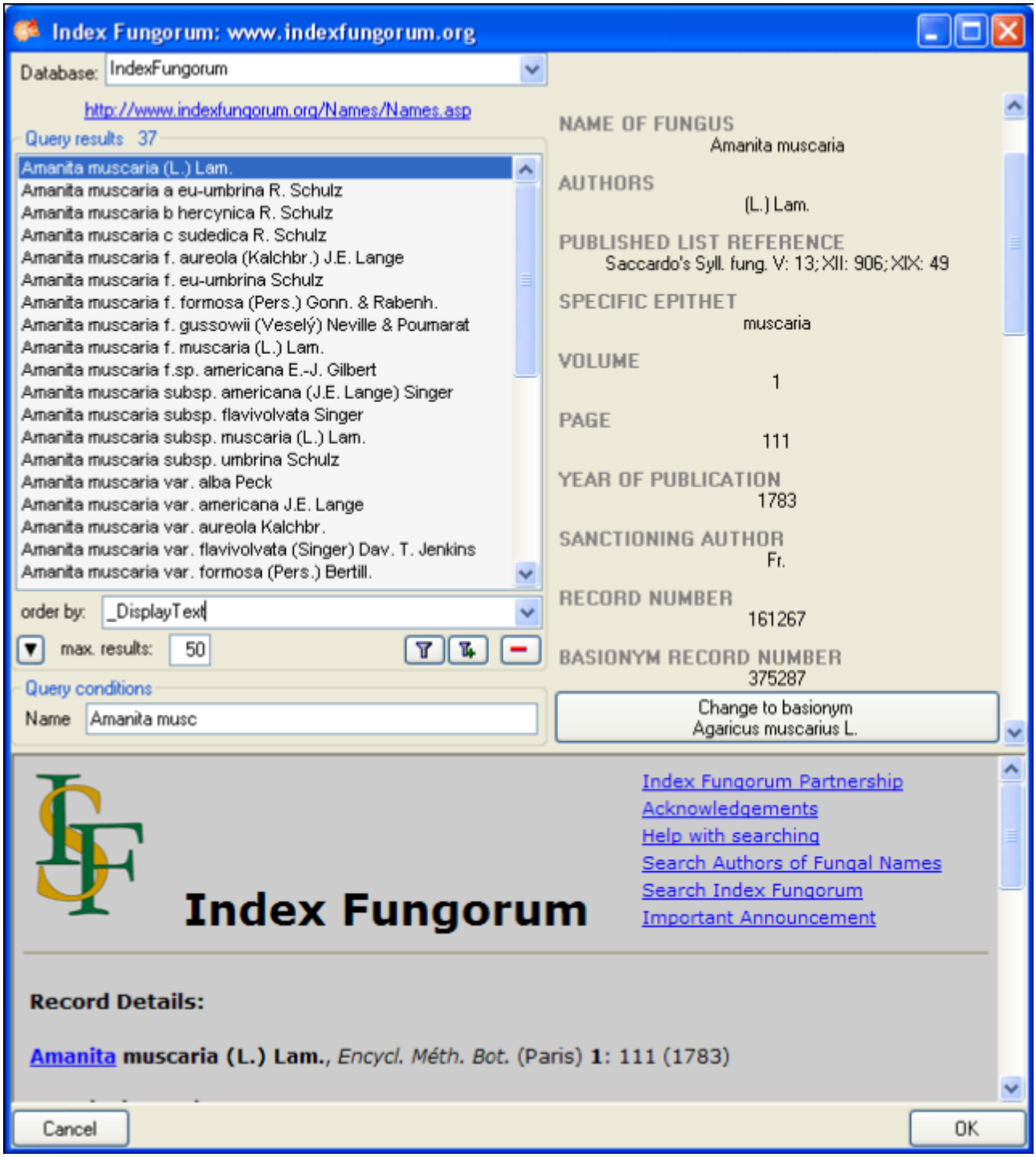
Index Fungorum

The Palaeontology Database

The Catalogue of Life

Index Fungorum - webservice

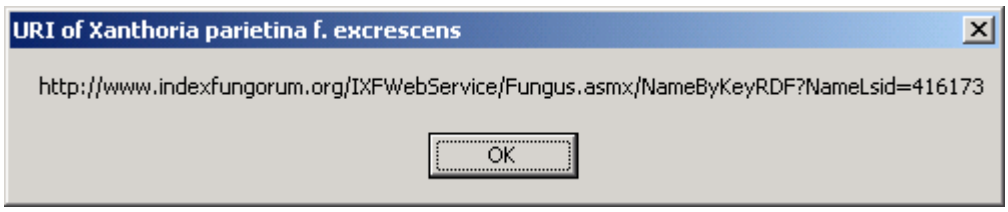
Some modules within the Diversity Workbench provide the possibility to link your data to an external webservice. For example DiversityTaxonNames gives you access to the taxonomic names of IndexFungorum. To establish a connection to this webservice, click on the button. A window will open where you can choose IndexFungorum from the Database list (see below).



Enter the query restriction for the name in the Name field in Query conditions. The maximal number of records you get can be set in the max. results field max. results: 50 (choose a low number if you have a slow connection to the internet). Then click on the search button to start the query. In the list of the left upper part the results of the query will be listed. In the

right part of the window additional information is shown as provided by the webservice. For certain entries buttons will appear, as e.g. shown above for the basionym and the current name of a scientific name. Click on these buttons if you want to change to one of these related datasets from the webservice. If available, the informations provided on the corresponding website is shown in the lower part. To take the link from the webservice into your database choose one of the entries and click OK. The entry will change as shown below.

If you double-click on the link area a window will open, providing you with the retrieval information of the webservice.

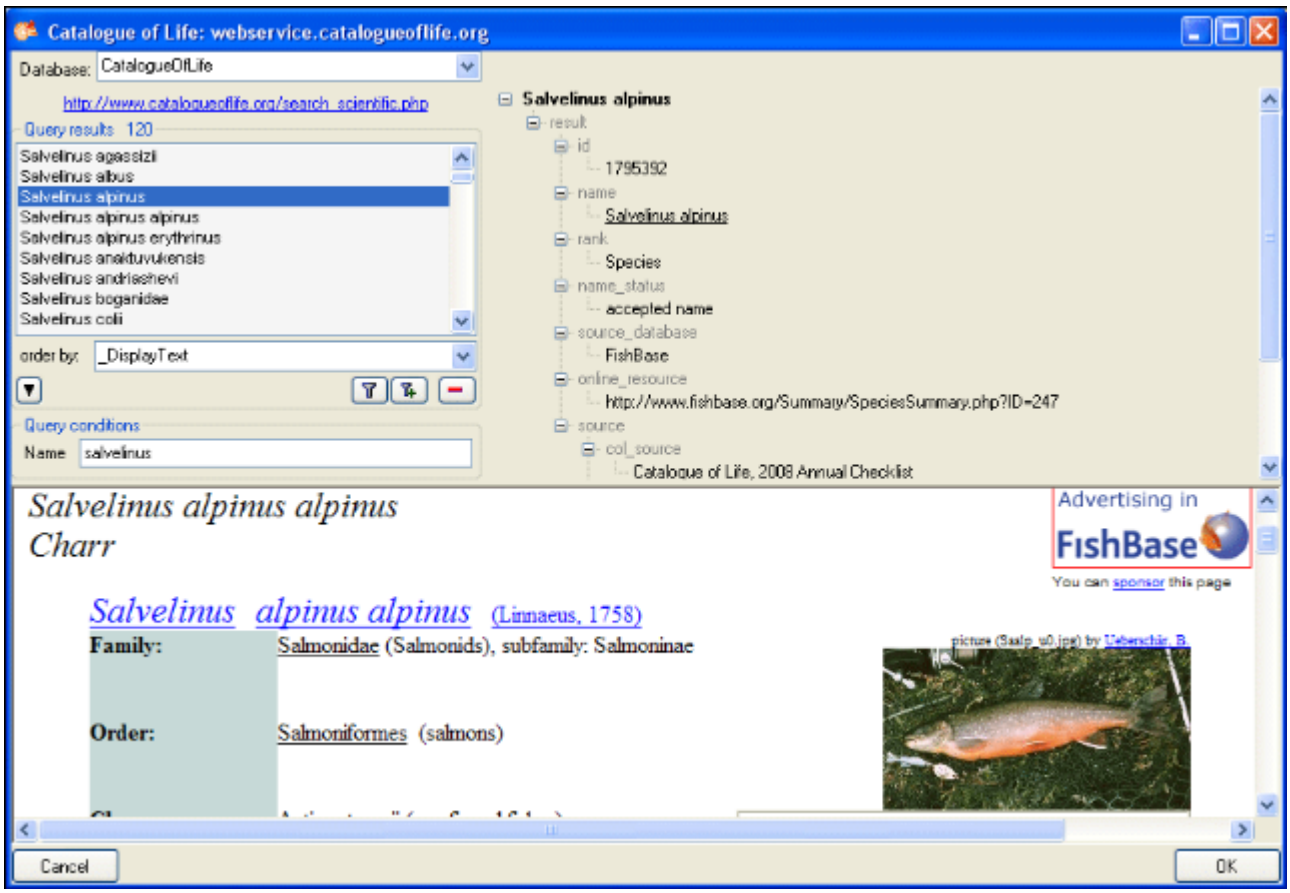


To get the whole information related to this entry as provided by the webservice, click on the button. A window will open as shown below where the informations of the webservice are listed, If available, the lower part will show the corresponding informations of a website.

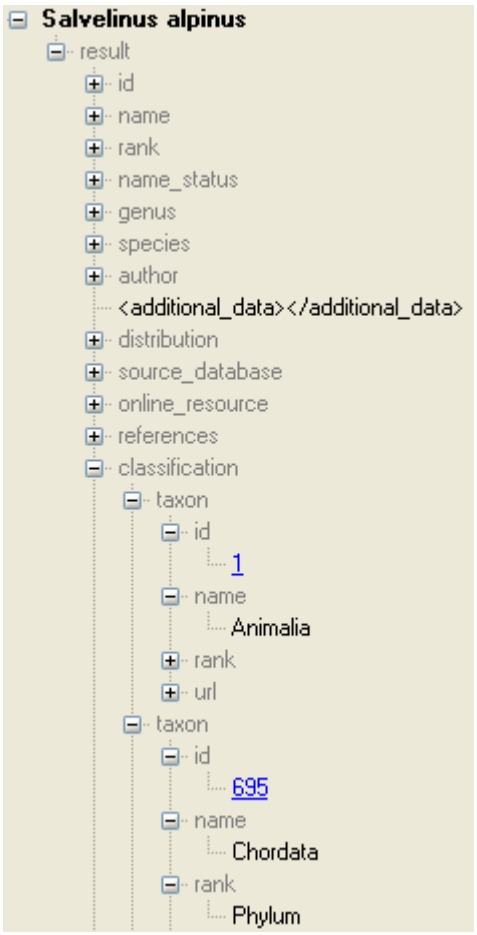
If you want to remove the link to the webservice, click on the the button. This will only remove the relation to the webservice, not the cached name.

Catalogue of Life - webservice

Diversity Workbench provide the possibility to link your data to an external webservice. The webservice provided by the Catalogue of Life is possible through the module DiversityTaxonNames. To establish a connection to this external webservice, click on the button. A window will open where you can choose this webservice (see below).

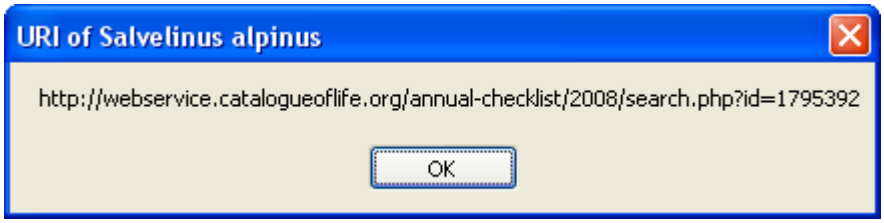


In the field **Name** in Query conditions enter you search string and click on the button to start the query. In the list of the left upper part the results of the query will be listed. In the right part of the window additional information is shown as provided by the webservice. If available, the lower part of the window will show the webpage of the related information.

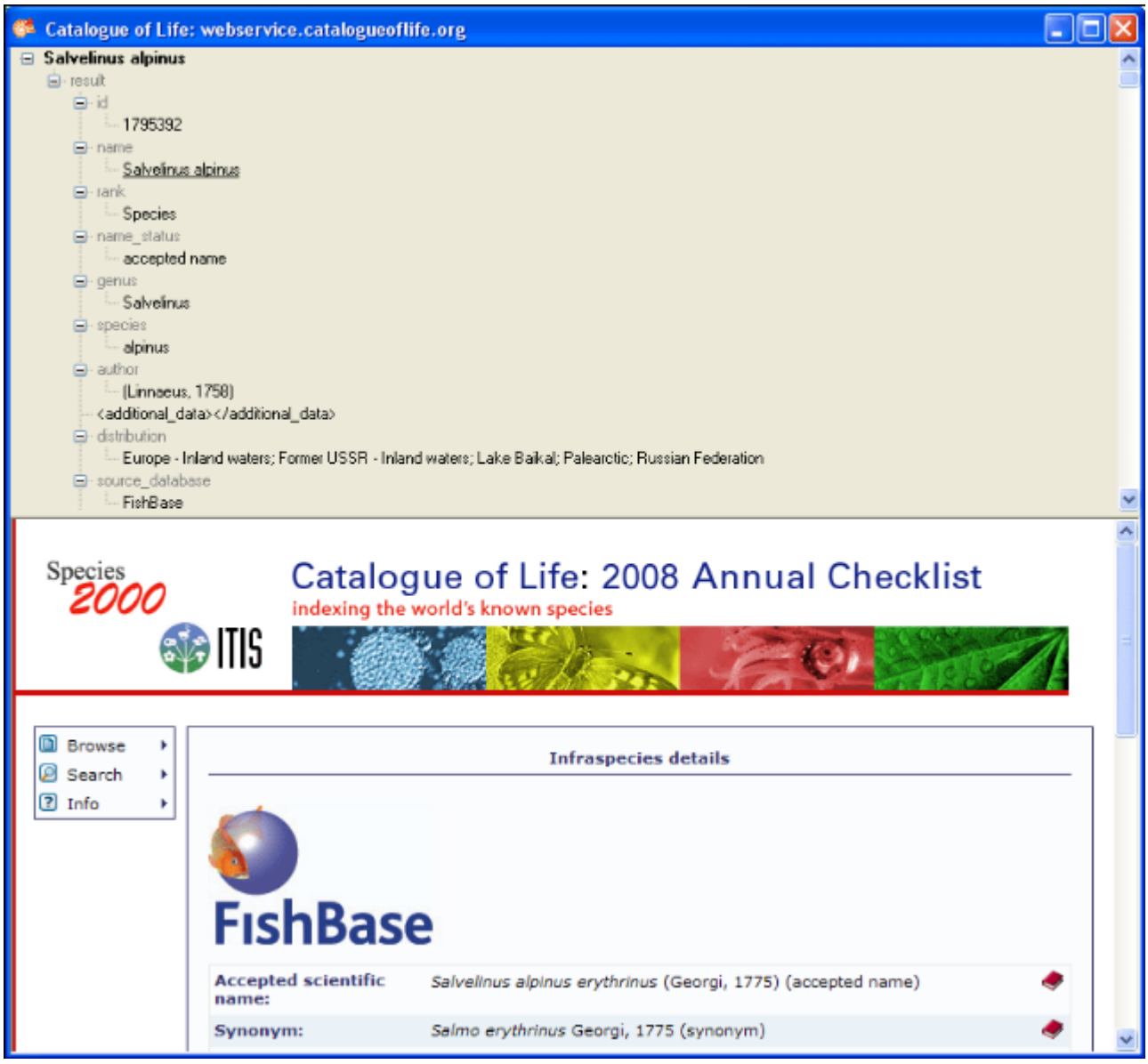


Higher taxa and for synonyms the accepted name will be shown as e.g. shown above. To inspect one of these entries, click on the linked entry of the ID - in the example above id: 695 . To take the link from the webservice into your database choose one of the entries in the list and click OK. The entry will change as shown below.

If you double-click on the link area a window will open, providing you with the retrieval information of the webservice.



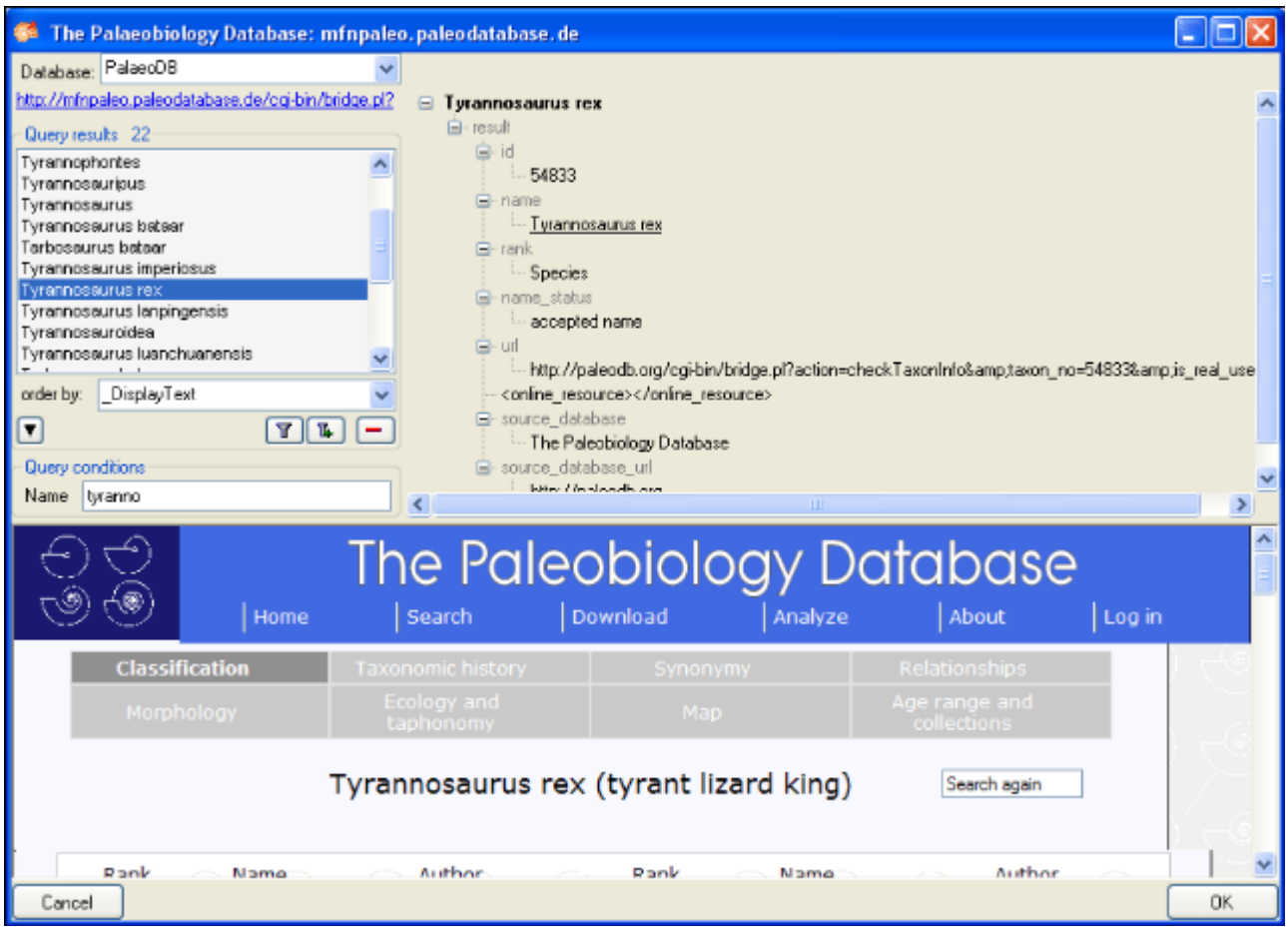
To get the information related to an entry as provided by the webservice, click on the button. A window will open as shown below where the informations of the webservice are listed in the upper part. If available, additional informations provided on a corresponding website will be shown in the lower part.



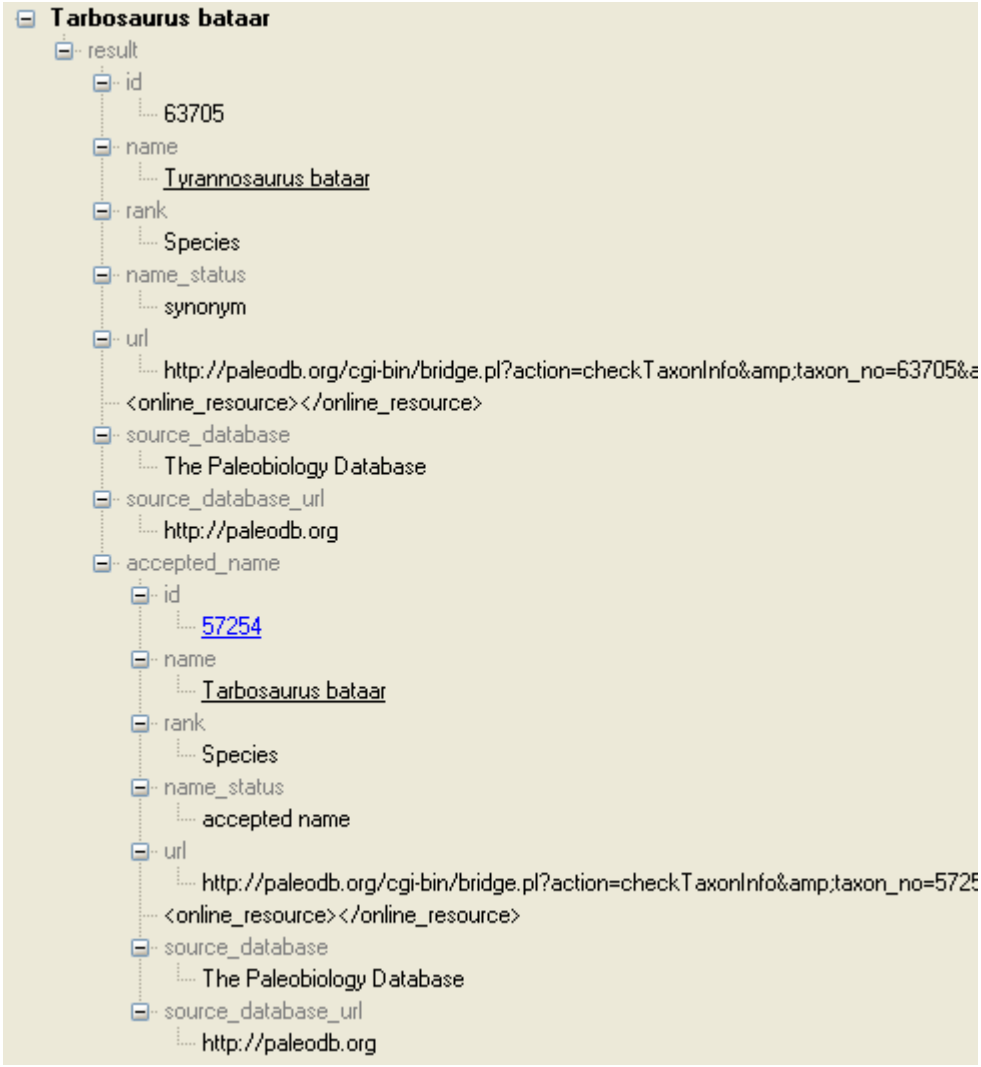
If you want to remove the link to the webservice, click on the ✕ button. This will only remove the relation to the webservice, not the cached name.

The Palaeontolgy Database - webservice

Diversity Workbench provide the possibility to link your data to an external webservice. The webservice provided by the Palaeontology Database is possible through the module DiversityTaxonNames. To establish a connection to this external webservice, click on the button. A window will open where you can choose this webservice (see below).



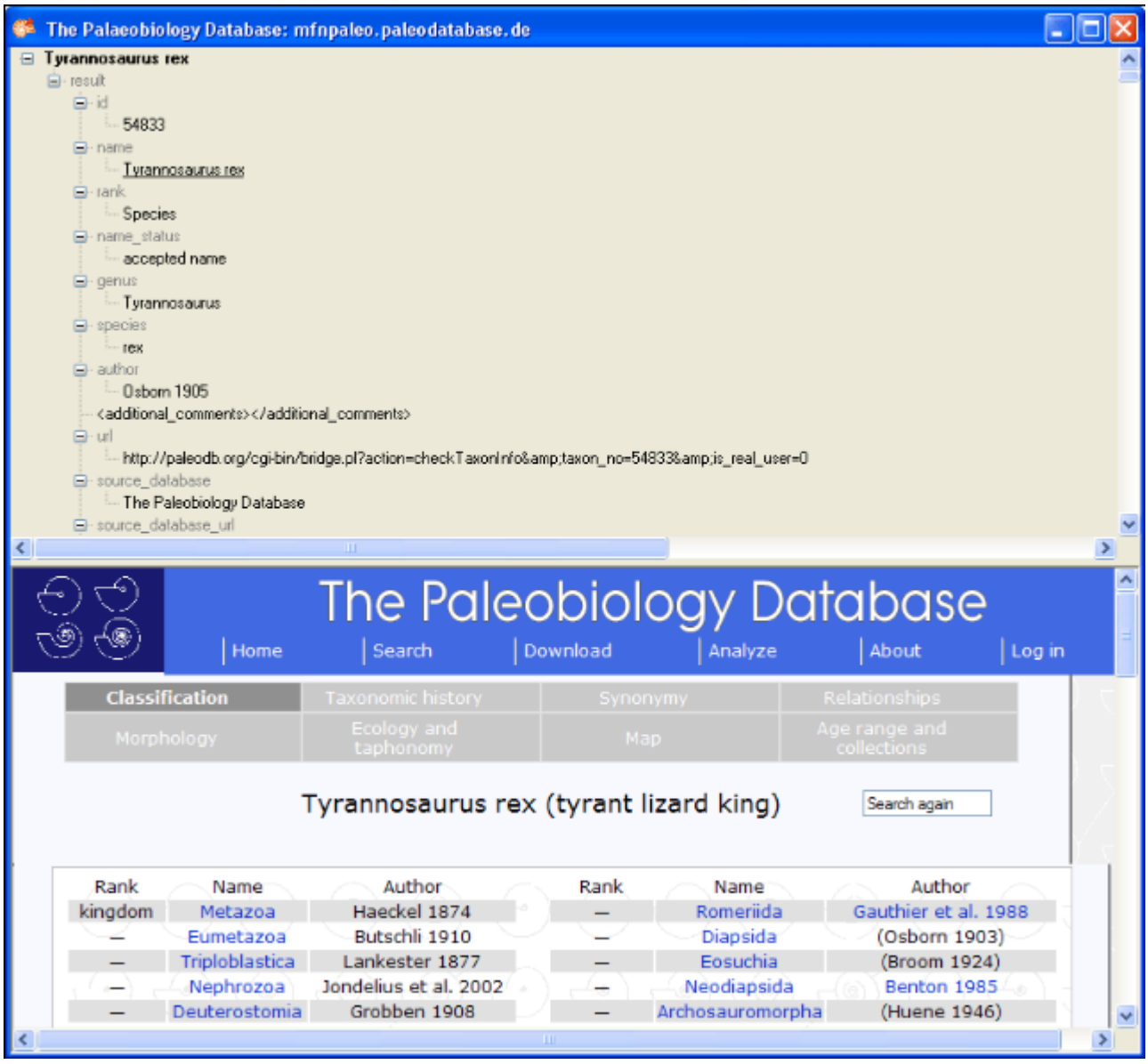
In the field **Name** in Query conditions enter you search string and click on the button to start the query. In the list of the left upper part the results of the query will be listed. In the right part of the window additional information is shown as provided by the webservice. The lower part of the window will show the webpage of the related information.



For synonyms, the accepted name will be shown as well as e.g. shown above. To change to the accepted name, click on the linked entry of the ID - in the example above id: 57254. To take the link from the webservice into your database choose one of the entries in the list and click OK. The entry will change as shown below.

If you double-click on the link area a window will open, providing you with the retrieval information of the webservice.

To get the information related to an entry as provided by the webservice, click on the button. A window will open as shown below where the informations of the webservice are listed in the upper part. If available, additional informations provided on a corresponding website will be shown in the lower part.



If you want to remove the link to the webservice, click on the ✕ button. This will only remove the relation to the webservice, not the cached name.