



R Guide

Version: 3.3.0
Date: 17.09.2019
Status: Valid
Author: A. Hueni, Remote Sensing Laboratories, University of Zurich
File: \SPECCHIO_R_Guide.docx
Pages: 16

Classification:
Distribution: SPECCHIO Users



History

Version	Date	Author	Remark
3.1.0	04.07.2014	A. Hueni	First version for SPECCHIO V3.1
3.3.0	17.09.2019	A. Hueni	Updates for account configuration

Table of Contents

1	Introduction	4
2	Installation and Configuration	5
2.1	Setup.....	5
2.1.1	Definition of the Java Classpath	5
2.1.2	Account Configuration.....	5
3	Connecting to the SPECCHIO Database.....	6
4	Interaction with SPECCHIO	6
4.1	Preparing Queries.....	6
4.1.1.1	Using the Data Browser to form Query Conditions	7
4.1.1.2	Using the Query Builder to form Query Conditions.....	7
4.2	Retrieving and Plotting Spectral Data	8
4.3	Retrieving Metadata.....	10
4.4	Updating Metadata.....	10
4.5	Inserting new Metadata.....	10
5	SPECCHIO API	10
5.1	SPECCHIOClient.....	10
5.1.1	getSpaces.....	10
5.1.2	getMetaparameterValues.....	11
5.1.3	updateEavMetadata.....	11
6	Examples	11
6.1	Function to get Spectral Data from SPECCHIO	11
6.2	Get Metadata of Spectra.....	12
6.3	Generating Time Series	14
7	Building Graphical User Interfaces	15
8	References.....	16

1 Introduction

The R environment is a popular tool in research and science. This integration of SPECCHIO with R bases on the rJava library¹, allowing the use of Java classes within R code.

The code introduced in this guide has been tested with version rJava_0.9-4.

The integration has been tested using RStudio; using other R environments are at the end-users own risk.

This guide is targeted at SPECCHIO Version 3.1 or higher.

¹ <http://cran.r-project.org/web/packages/rJava/index.html>

2 Installation and Configuration

2.1 Setup

2.1.1 Definition of the Java Classpath

The java classpath of the rJava package needs to be updated to include the SPECCHIO classes. This can be done dynamically within each R script using the SPECCHIO functionality by specifying the path to the `specchio_client.jar` file.

Under MacOSX for the installation compiled as MacOS application, the default path would be:

```
.jaddClassPath(path="/Applications/SPECCHIO/SPECCHIO.app/Contents/Java/specchio-client.jar")
```

For the cross-platform installation, the path would depend on your installation path:

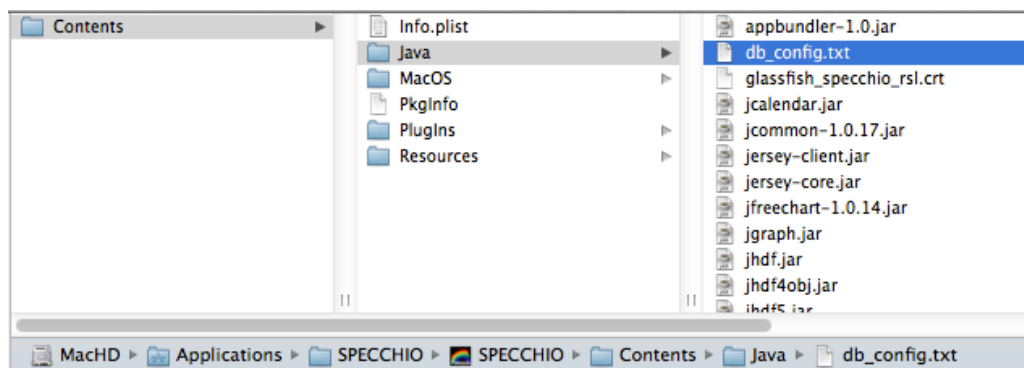
```
.jaddClassPath(path="<installation-folder>/specchio-client.jar")
```

2.1.2 Account Configuration

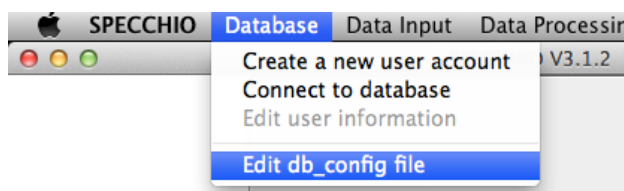
Using SPECCHIO from R automatically uses your existing account configurations, either stored in the Java Preferences on the operating system, or stored within a file named `db_config.txt`.

The `db_config.txt` file is found in your installation path of SPECCHIO in the same directory as the `specchio-client.jar` file.

E.g. under MacOSX:



To see the content of the `db_config.txt` start the SPECCHIO Java application and select the *Database* and *Edit db_config file* menu item:



This will open the `db_config.txt` file in a text editor.

The order of the connections within this configuration file will be the identical to their order within a list of SPECCHIOServerDescriptor objects. The server descriptors are returned as a Java ArrayList and are therefore indexed from zero onwards.

The code to select a connection from the ones stored in the configuration file and connect to SPECCHIO is covered below (see Section 3).

3 Connecting to the SPECCHIO Database

Use the SPECCHIOClientFactory class to open a connection to a SPECCHIO database instance:

```
# connect to server
specchio_cf <- J("ch/specchio/client/SPECCHIOClientFactory")$getInstance()
```

This call reads the known database connections from the SPECCHIO db_config.txt file. All known server connections can be returned as an ArrayList object by calling the getAllServerDescriptors method;

```
descList <- specchio_cf$getAllServerDescriptors()
```

Connect to any entry in the descriptor list by calling the createClient method, parameterized with an entry from the descriptor list. E.g. to connect to the first entry in the list, get the zero-th entry (Java lists are indexed from zero onwards):

```
# connect using first connection description
desc <- descList$get(as.integer(0))
specchio_client <- specchio_cf$createClient(desc)
```

The SPECCHIOClient class uses the Singleton pattern (Gamma, Helm et al. 1997). This implies that once the connection is set, all SPECCHIO objects connecting to the database automatically utilise this connection.

The specchio_client object provides all methods to interact with the SPECCHIO database.

To get the number of database connections stored in the db_config.txt file use:

```
descList$size()
```

To list a specific entry, use:

```
descList$get(as.integer(2))
```

```
[1] "Java-Object{https://AHueni@v473.vanager.de:443/specchio_service@jdbc/specchio_prod}"
```

4 Interaction with SPECCHIO

4.1 Preparing Queries

Queries conditions are used to select spectral data from the database. These can be defined manually, but the easiest way is to generate queries using the SPECCHIO Data Browser or Query Builder and use these in R.

These automatically created queries are a good starting point to programmatically create query conditions, e.g. select data within a loop with changing query conditions.

4.1.1.1 Using the Data Browser to form Query Conditions

Start the regular SPECCHIO Java Application, open the Data Browser and select the required data. Then click the menu button over the 'Matching Spectra' text area and select 'Copy R-ready query to clipboard'.

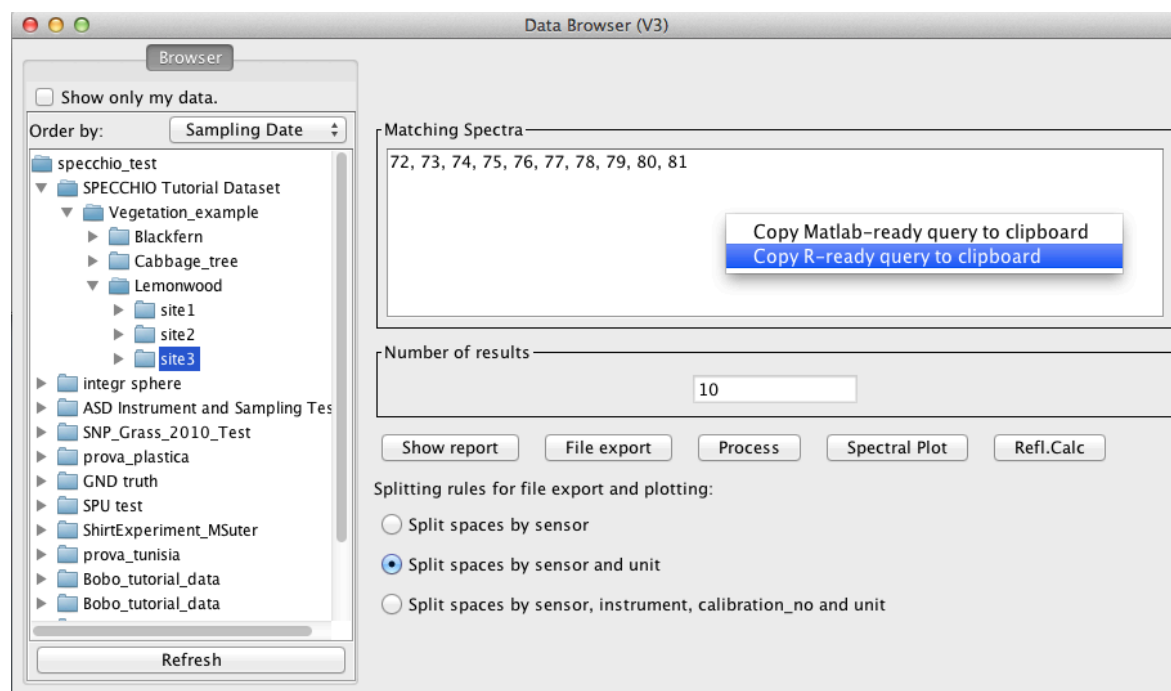


Figure 1: R query pop-up menu in the Data Browser

4.1.1.2 Using the Query Builder to form Query Conditions

Start the regular SPECCHIO Java Application, open the Query Builder and select the required data. Then click the menu button over the 'Matching Spectra' text area and select 'Copy R-ready query to clipboard'.

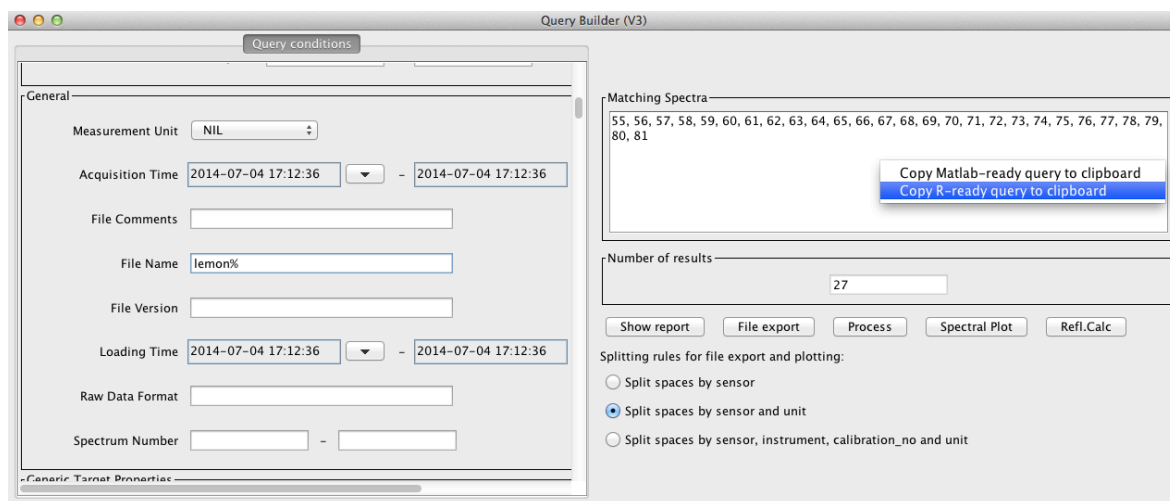


Figure 2: R query pop-up menu in the Query Builder

4.2 Retrieving and Plotting Spectral Data

Data is retrieved from the database by the following steps:

Paste the query conditions (see 4.1 on how to create queries) in your R file.

This will be similar to:

```
query <- .jnew("ch/specchio/queries/Query", "spectrum")
query$setQueryType(query$SELECT_QUERY)

query$addColumn("spectrum_id")

cond <- .jnew("ch/specchio/queries/QueryConditionObject", "spectrum", "spec-
trum_id");
id_array <- c(72,73,74,75,76,77,78,79,80,81)
ids_list <- .jnew("java.util.ArrayList")
for (i in 1:length(id_array)) { ids_list$add(.jnew("java.lang.Integer",
as.integer(id_array[i]))) }
cond$setValue(ids_list);
cond$setOperator("in");
query$add_condition(cond);

ids <- specchio_client$getSpectrumIdsMatchingQuery(query);
```

This code returns the spectrum ids that identify the spectra matching the query conditions.

Call the `getSpaces` method (set to split spaces by sensor only and order the matching entries by their acquisition time) (see 5.1.1):

```
spaces <- specchio_client$getSpaces(ids, TRUE, FALSE, "Acquisition Time")
```

The spaces object is an array of spaces created by the Spectral Space Factory, a service offered by the SPECCHIO server and essentially creating containers to hold spectra of different wavelength dimensions.

To figure out how many spaces were returned, use:


```
spaces$length
[1] 1
```

To get a particular space, use the standard R array syntax, e.g. get the first element of the space array:

```
space <- spaces[[1]] # first element of returned array
```

Spaces are just pre-configured containers to hold spectral vectors. To actually populate a space, use the `loadSpace` method, supplying the space required loading as an argument:

```
space <- specchio_client$loadSpace(space);
```

Once a space is loaded use the method `getVectorsAsArray` to get the spectra held by the space as array and the method `getAverageWavelengths` to get the centre wavelengths (i.e. average of the spectral response function per sensor band):

```
# get spectra and wvls
vectors <- space$getVectorsAsArray()
wvls <- space$getAverageWavelengths()
```

To convert the Java array holding the vectors as R matrix, use

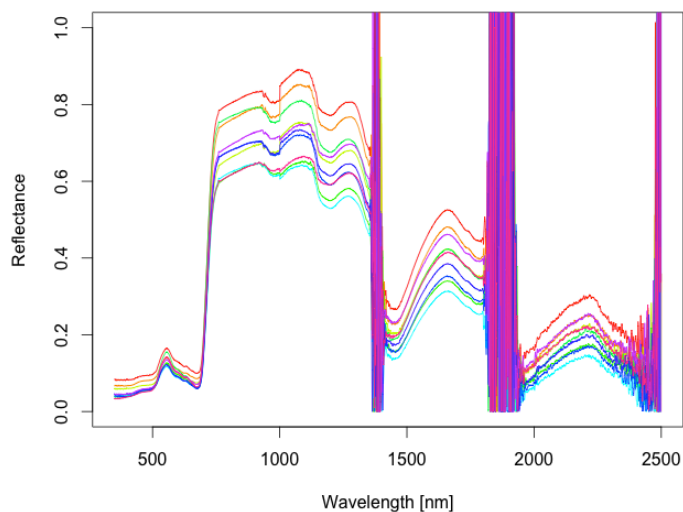
```
# convert java array to R matrix
r_matrix <- .jvalArray(vectors, simplify=TRUE)
```

The space holds information about the measurement unit as well, hence, all spectra can be plotted and the plot annotated as follows:

```
plot(r_matrix[1,]~wvls,type="n", ylim=c(0, 1), xlab="Wavelength [nm]" ,
ylab=space$getMeasurementUnit()$getUnitName()) # create empty plot
```

```
col = rainbow(space$getNumberOfDataPoints())
```

```
for (i in 1:space$getNumberOfDataPoints() ) {
  lines(r_matrix[i,]~wvls, col=col[i])
}
```



4.3 Retrieving Metadata

Metadata are easily retrieved for a selection of ids by calling the `getMetaparameterValues` method of the `SPECCHIOClient` (5.1.2) class, e.g.:

```
latitudes <- specchio_client$getMetaparameterValues(ids, "Latitude")
```

If data are numerical, they are returned as a special list object of the class `MatlabAdaptedArrayList`, allowing an easy conversion of the data held by the returned list into a Matlab or R array via the method `get_as_double_array()`.

Otherwise, e.g. for string data types, values are returned in a Java `ArrayList`.

For an example on how to read metadata from SPECCHIO see the example provided in section 6.2.

4.4 Updating Metadata

Existing metadata can be updated programmatically by using the `updateEavMetadata` method of the `SPECCHIOClient` () class.

4.5 Inserting new Metadata

New metadata can be inserted programmatically by using the `updateEavMetadata` method of the `SPECCHIOClient` () class.

The following example inserts a new Boolean metaparameter to flag a selection of dark current spectra.

```
% get attribute object for the metadata attribute 'DC Flag'
flag_attribute = specchio_client.getAttributesNameHash().get('DC Flag');

% insert flag by creating a new metaparameter, setting the value and updating the
% metadata of selected spectra
e = MetaParameter.newInstance(flag_attribute);
e.setValue(1);

specchio_client.updateEavMetadata(e, ids);
```

5 SPECCHIO API

5.1 SPECCHIOClient

5.1.1 getSpaces

Get the space objects for a set of spectrum identifiers.

```
* @param ids                the spectrum identifiers
* @param split_spaces_by_sensor  boolean
* @param split_spaces_by_sensor_and_unit  boolean
```

```
* @param order_by           the field to order by
```

The Boolean flags allow to configure the space building as in the SPECCHIO Data Browser or Query Builder Interfaces.

5.1.2 getMetaparameterValues

Get values for spectrum ids and EAV attribute.

```
*
*
* @param ids           spectrum ids
* @param attribute     attribute name
*
* @return list of values, or null if the field does not exist
```

Note: There is currently no check implemented to ascertain that all spectra defined by the ids do have such an attribute defined; it can thus happen that the number of values returned is lower than the number of spectra in the list!

5.1.3 updateEavMetadata

Updates or inserts a metaparameter for a specified list of spectra.

```
* @param mp           the meta-parameter to update
* @param spectrum_ids the identifiers for which to update the parameter
*
* @return the identifier of the inserted metadata
```

```
updateEavMetadata(MetaParameter mp, ArrayList<Integer> spectrum_ids)
```

```
* @param mp           the meta-parameter to update
* @param spectrum_ids the identifiers for which to update the parameter
* @param mp_old       the old meta-parameter
*
* @return the identifier of the inserted metadata
*/
public int updateEavMetadata(MetaParameter mp, ArrayList<Integer> spectrum_ids,
MetaParameter mp_old)
```

6 Examples

6.1 Function to get Spectral Data from SPECCHIO

This script is the complete example of code introduced in sections **Error! Reference source not found.** - 4.2.

```
library(rJava)
.jinit()
```


8 References

Gamma, E., R. Helm, et al. (1997). Design Patterns. Elements of Reusable Object-Oriented Software. Reading, MA, Addison Wesley Longman Inc.