

Manual for Elchemea Analytical

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January 15, 2013

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Chapter 1

Introduction to Elchemea Analytical

The Department of Energy Conversion and Storage at Technical University of Denmark Impedance visualisation and analysis control software system (Elchemea Analytical) is a generalised visualisation / fitting software package for visualising impedances data. The Elchemea Analytical system is based on Perl and Apache and all graphics/fitting is done using Gnuplot®.

The main features of Elchemea Analytical are listed below:

- Simulation of impedance spectra using a wide variety of discrete impedance elements (R,C,L,Q,RQ,RC,W,O,G,T) as well as parallel and/or series connections of those.
- Fitting of impedance models to measured impedance data in a wide frequency range (from 10^{-100} to 10^{100} Hz).
- Easy determination of start parameters for a wide range of the predefined impedance elements (RC,RQ,O,G,T) using the 'find-values' built in algorithms.
- Easy integration to Elchemea© and Risoe Fuel cells and solid stage chemistry division Fuel Cell Control system© impedance acquisition and test system control software packages.
- Uses only open source software (OSS).

The first part of this documentation is an overview of the user interface (section 3) mainly intended for new users of the system. The second part (chapter 6) is mainly intended for more advanced users and system administrators as it contains information regarding configuration. It is assumed that any administrator has a fairly advanced knowledge of Unix system administration and Perl programming.

Chapter 9 contains the documentation for the different perl module supplied by the Department of Energy Conversion and Storage at Technical University of Denmark Impedance visualisation and analysis control software.

Chapter 2

License

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Chapter 3

User interface

The Department of Energy Conversion and Storage at Technical University of Denmark Impedance visualisation and analysis control software system is based on the Apache web server software (Open Source Software, OSS). In figure 3.1 the main page is shown as an example of the web pages.

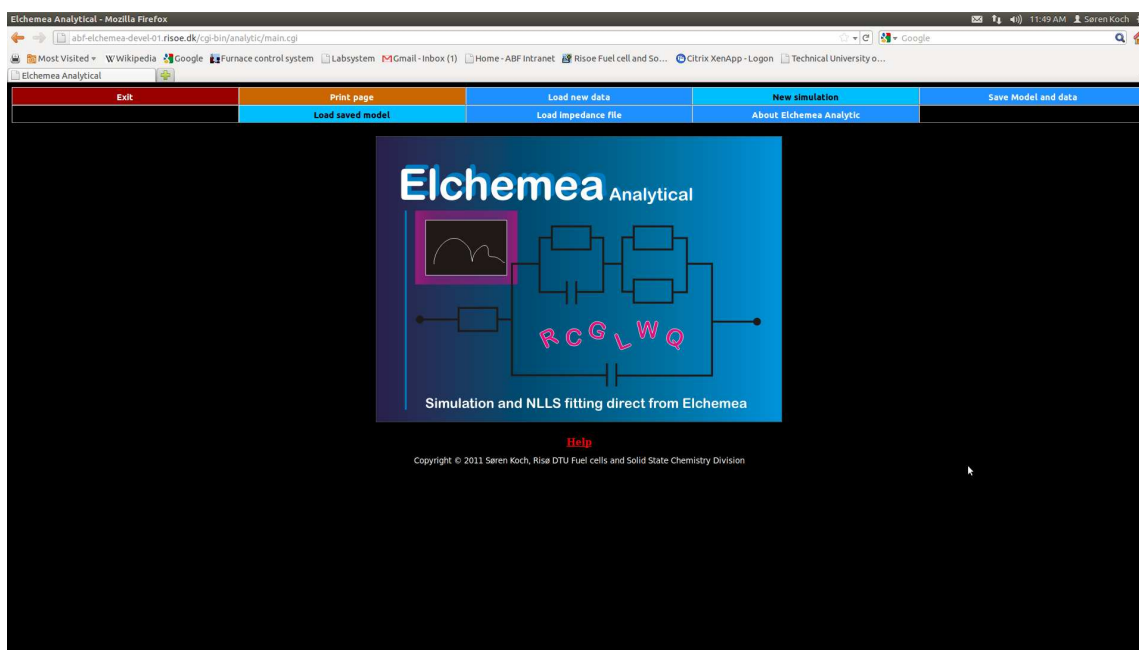


Figure 3.1: The main page.

3.1 Simulation

In order to simulate spectra, press the 'New simulation' button, and then build your impedance model element by element by using the 'Add element' button. In general, all elements will be placed in series, except if they are part of an explicit parallel or serial connection (by using the 'Par' or 'Ser' elements at the bottom of the element list). Figure 3.2 shows an example of a simple series connection of three impedance elements,

a Gerisher element (G) a RC element (parallel connection of a resistor and a capacitor) and an inductor (L).

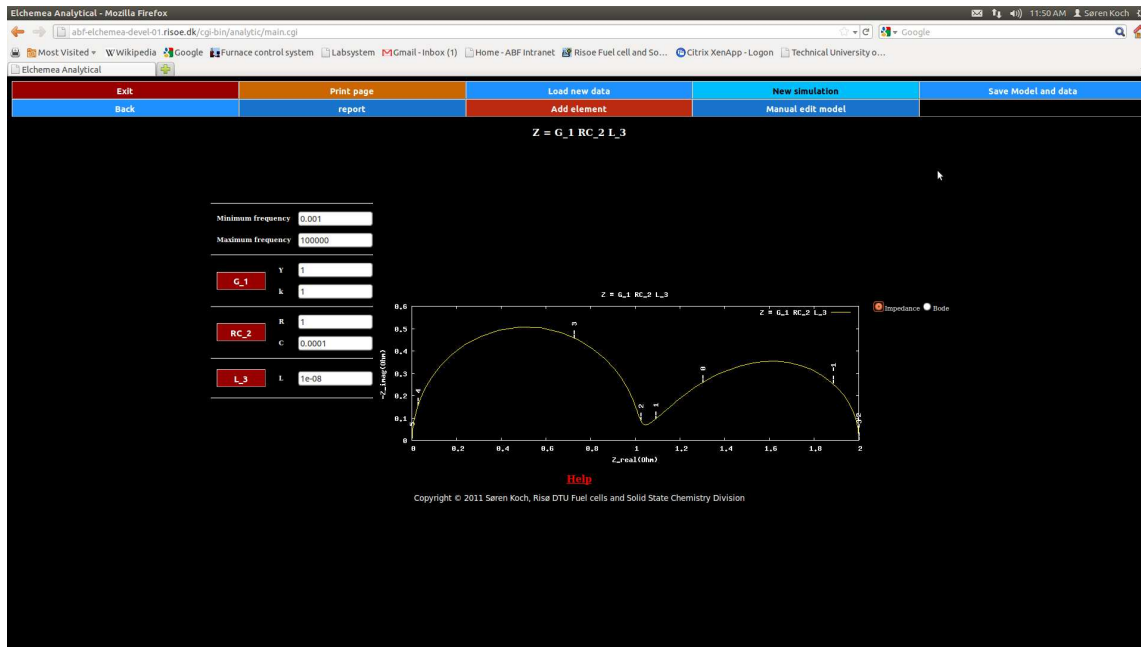


Figure 3.2: An example impedance simulation using simple series connection of three impedance elements..

If custom series and / or parallel connection elements are used, the model description on the left of the page looks slightly different than the one shown in figure 3.2. Figure 3.3 shows an example of how this may look. The way to read the complex models presented in this way is as follows:

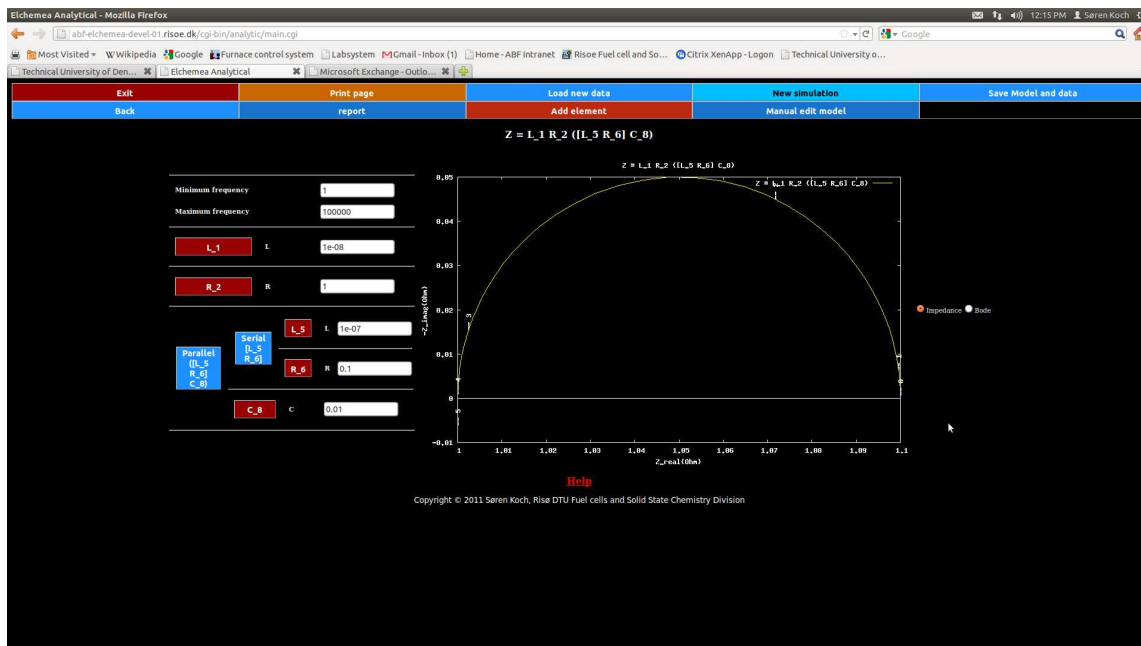


Figure 3.3: An example impedance simulation using parallel and serial connections of discrete elements.

The first two red buttons (labeled 'L_1' and 'R_2') are simple impedance elements connected in series (as normal) and in series with the third element represented by the leftermost blue button (labeled 'Parallel ([L_5 R_6] C_8)'). This button (labeled 'Parallel ([L_5 R_6] C_8)') indicates a parallel connection of all elements between the lines above and below this button. In this case it is a parallel connection of two elements; an capacitor ('C_8') and a series connection (labeled 'Serial [L_5 R_6]'). The series connection (represented by the second blue button) contains all elements to the right of it which is between the white lines above and below the blue button (in this case 'L_5' and 'R_6').

Thus the complete model is $Z = L_1 + R_2 + \text{Par}(\text{Ser}(L_5 + R_6), C_8)$.

In this way it is possible to build arbitrary complex impedance models, either for simulation purposes or for fitting to actual measured spectra.

If an element is to be deleted, simply press the red button designating the element in question. If deleting elements results in an empty container (parallel or serial), then the container itself is also deleted.

It is also possible to manually edit the impedance model. To do this press the 'Manual edit model' button, this brings up a small editor where the model is displayed. The model consists of a number of lines, each designating a single element (an exception to this is the container elements, which takes up 2 lines). The container elements start with a line with a single bracket. Angled brackets '[' for series connections and normal brackets '(' for parallel connections. This is similar to the nomenclature used in the old DOS program 'equivrt' by Dr. Bernard A. Boukamp. The corresponding line with the end bracket denotes the end of the container element. Note that containers may contain containers! Also note that if you are manually editing models, keep track of the brackets, as misaligned brackets may invalidate the model and / or result in an unwanted model!

3.2 Data analysis

In order to analyse measured impedance spectra, use one of the load options on the main page (figure 3.1). Figure 3.4 shows how a freshly loaded data file may look. If no prior model has been defined, the Elchemea Analytical generates a simple impedance model to start on. This consists of an inductor in series with a resistor and the values are determined from the data points in the highest frequency decade. The real value of the data point corresponding to the highest frequency determines the resistance and the inductance is determined by doing a Kramers Kröning transformation on the data points in the highest frequency decade (Algorithm and program supplied by Christopher Graves). Notice however that in case the Kramers Kröning method fails the imaginary value of the first data point (highest frequency) is used instead. This is to ensure that a value is always obtained, and this is known to happen for some combinations of Python and SE-Linux.

By using the radio button on the right it is possible to switch between Nyquist view and Bode view.

If some data points are to be excluded from the analysis, click on the data graph and then maneuver the pointer to the data point which is to be deleted and press the 'Delete'

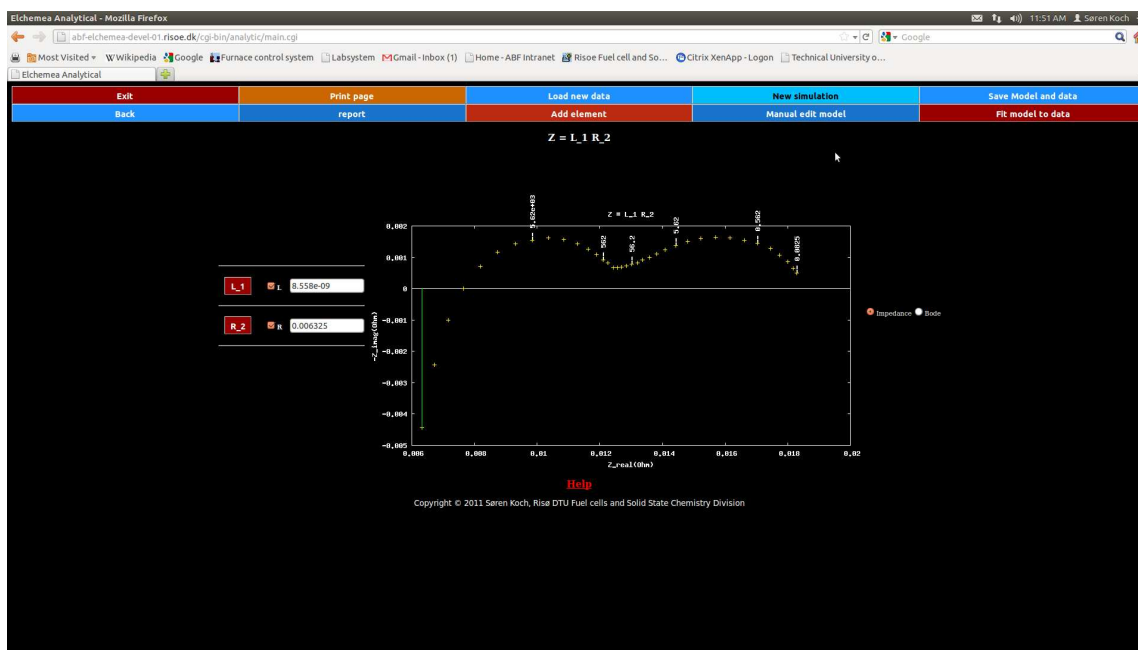


Figure 3.4: Example of an freshly loaded impedance spectrum.

button. Repeat this procedure until the data set has been sanitised from any erroneous data.

Caution: Do not delete data points merely because they do not fit the chosen model!

Only delete data points if they are known to be problematic (a single point lying far away from the rest is a good candidate, but not a sure sign)!

A special feature of the Elchemea Analytical is the ability to determine acceptable start values for parameters for some of the predefined complex impedance elements (RC, RQ, O, G and T). To use this, select the appropriate element to add, and then press the 'Find values' button, which will bring up an additional window like the one shown in figure 3.5.

Use the arrow buttons to move the frequency marker to the desired data points and select the right data points using the three blue buttons below ('Set maximum frequency', 'Set summit/bend frequency' and 'Set minimum frequency'). Once these three frequencies have been selected (which can be seen on the display as changes in the color and symbol of the data points), press the red 'Find values' button. This will cause the Elchemea Analytical system to try and find sensible start parameters for the selected data range for the element in question, and in case a fit converges the resulting impedance model will be shown as seen in figure 3.6. If you are satisfied, simply press the 'OK' button (which causes the find values window to close) and then the 'Add element' button, which will add the chosen element with the determined start parameters to the current model. It is possible to zoom using the X-range fields at the bottom (for instance if a small arc sits next to a large one, the small one may be extremely hard to determine). Simply change the minimum or maximum impedance to be displayed and the plot/select area will be updated accordingly.

Once a suitable model has been built, fit the model to the measured data by pressing

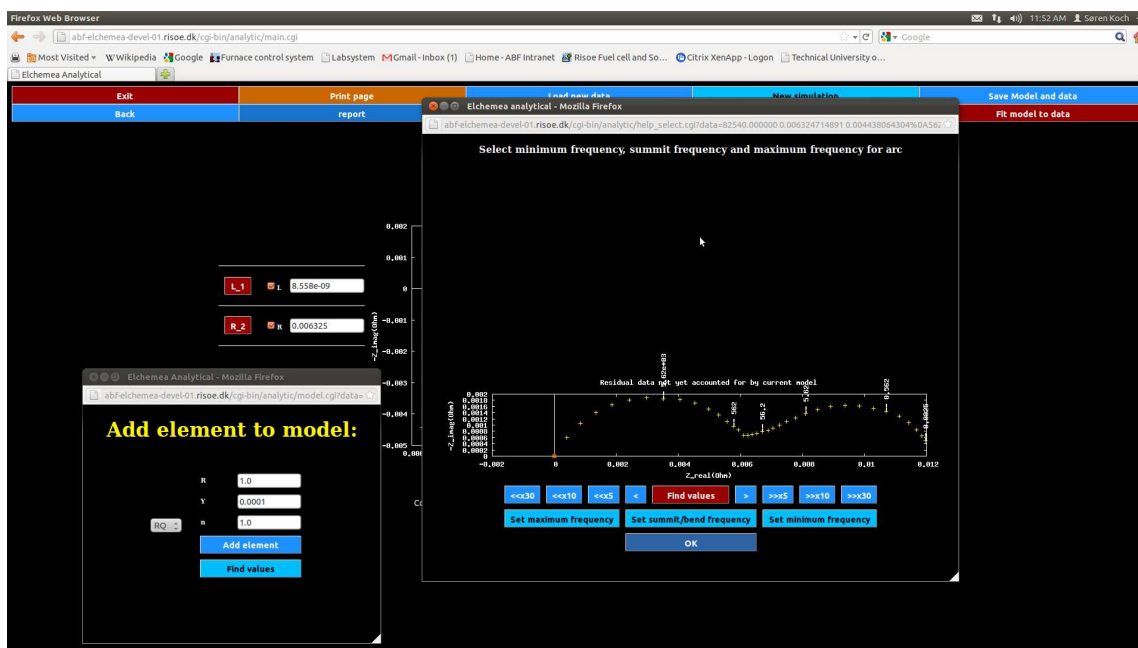


Figure 3.5: Adding an impedance element to a model using the 'find-values' system.

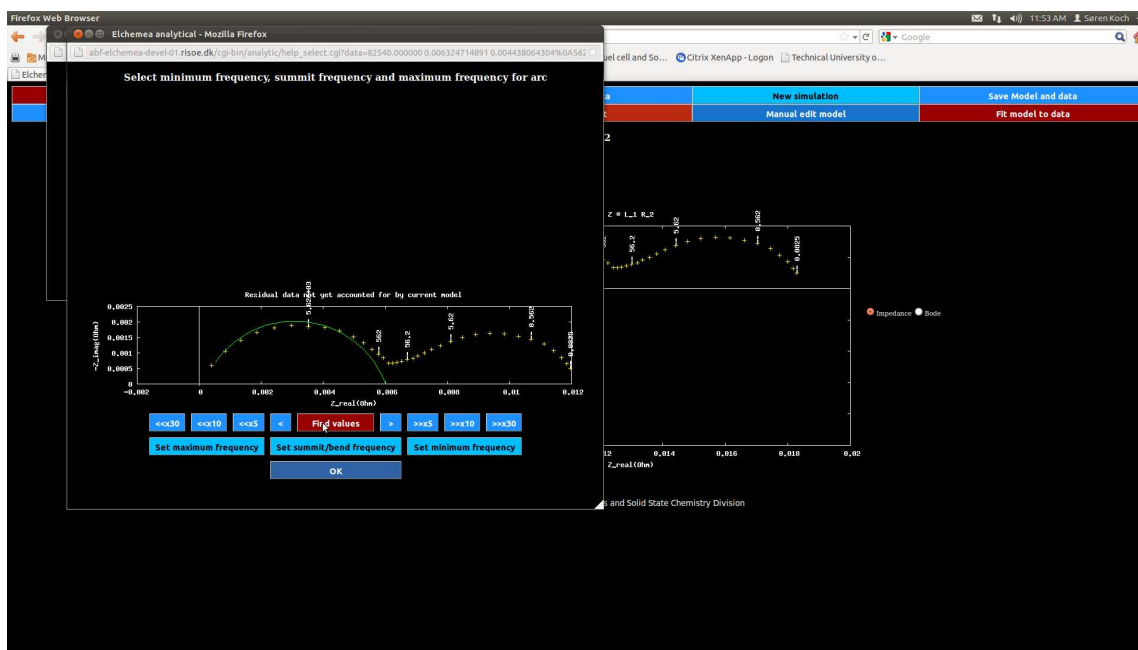


Figure 3.6: Finding the values for an RQ element using 'find-values' on a measured impedance spectrum.

the 'Fit model to data' button. This will cause the Elchemea Analytical to try and fit the model to the data, and in case the fit converges, the resultant parameter values will be displayed and the system will ask if the values should be copied to the model. If the resulting values are sensible, press 'Yes' and the resulting fit parameters will be displayed in the model section as well as appended to the fit result table. This table contains a line for each accepted fit and includes all model parameters as well as the mean and maximum error for the fit in question. Thus if more than one file is fitted (or a single data set is

fitted to more than one model), it is possible to get a list of all the fitted data afterwards for later comparison / further data analysis. To access the fit result table, press the 'Show fit table' below the graph. This brings up a new window with a tab delimited table which can be copied / saved to a local file as appropriate. Figure 3.7 shows an example of a fit result.



Figure 3.7: Example result of a fit of a model to an actually measured impedance spectrum.

It is possible to exclude some parameters from the fitting (in effect locking them to the current value) by unclicking the checkbox associated with the parameter in question. This is useful if external information indicates that a specific impedance parameter must have a specific value. After the fit has been run, press the 'Report' button to get a pdf report of the fit in question. This report will include parameter values, statistics regarding accuracy of the fit, parameter correlation as well as Nyquist, Bode and error plots of the model and data.

It is known that some versions of AdobeReader® in some cases has problems displaying the pdf documents created by Elchemea Analytical, however the fast open source pdf viewer Ghosview® (<http://pages.cs.wisc.edu/~ghost/>) is available for free for a wide range of operating systems.

3.3 Saving and reloading models

Once a model has been created it is possible to save the model (and any associated data). To do so, press the 'Save model and data' button, which will open a save dialog asking where to place the save data.

In order to load a saved dataset/model, press the 'Load saved model' button on the main page and select the data file to load (The default extension will be .ea).

It is also possible to save just the model, if this is to be desired, simply open the 'manual edit model' page and copy the model to the clipboard and then save it in a local text file. To restore a model thus saved, open a new simulation or data file and press the 'manual edit model' button and then paste the saved model instead of the present model. Finish by pressing 'ok'.

3.4 Hints on fitting

Fitting impedance data is as much an art as it is science. The most important part of data modeling / fitting is to select the most appropriate model, and in this respect Ockhams Razor is an extremely important tool! Do not select a more complex model than necessary based on the available information as it is always possible to get a good fit if enough impedance elements are added to the model.

Due to the mathematics in some of the impedance elements distributed with Elchemea Analytical, sometimes the fitting will not terminate or crash (indicated by either an error message or simply that no response is received from the fitting attempt). Especially the Gerisher, Finite length Warburg and bounded Warburg elements are prone to this as they include one or more of the hyperbolic trigonometric functions, and in some cases the fitting algorithm will select values which results in a singularity (division by zero or infinite) and the fitting algorithm fails. When this happens, try to slightly change the start parameters and see if it is possible to avoid the singularity.

3.5 Batchfitting

It is possible to do batch process fitting without using the GUI. To do this use the command line program *batchfit.pl* (usually located in */usr/local/bin/analytic/*). If the program is called without any arguments a description of how to use it is output. In the *batchtest* directory found in the Elchemea Analytic distribution directory there is an example of how to use it. Simply run the *run_batchfit_test.bash* program to test with the files in that directory.

Chapter 4

Plotting multiple files

It is also possible to plot multiple impedance file in the same plot by using the multiplot module of Elchemea Analytical. To access this module, navigate to the front page and press the 'Plot multiple files' button, which brings the multiplot page up as shown in figure 4.1.

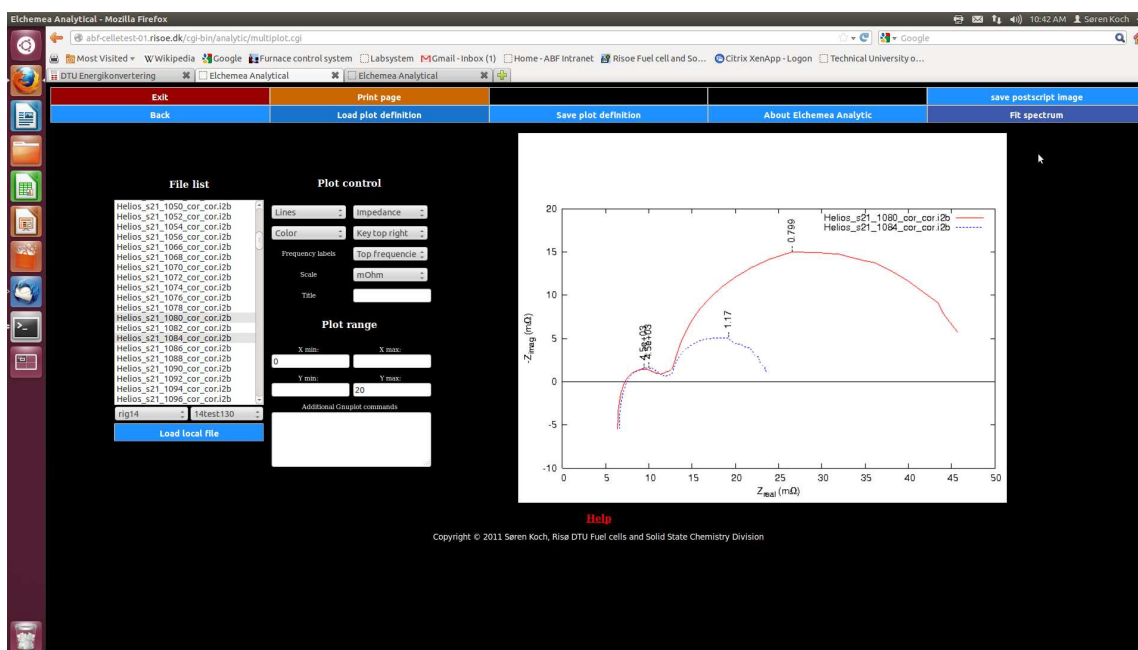


Figure 4.1: Multiplot page showing an example of 2 impedance spectra plotted on the same graph.

The left part is the file control section. It contains a multi select box showing both the available and selected files. To plot files load the impedance files from your local system by pressing the 'load local file' or select username, session number and files (in case the Elchemea@or RFCcontrol@software is installed on the server along with Elchemea Analytic). Note that the user/rig name and test/session select boxes are only available as indicated above. If files are loaded from a local resource, it is automatically selected for plot. To deselect/deselect a file press 'Ctrl' and click on the file to select/deselect. If the order of the legend keys are to be changed, move the file names up and down in the

file list as appropriate as Elchemea Analytics always uses the files/legends in the order they are shown in the file list. To move a file, select it and use the 'u' and 'd' keys to move the file up and down in the list (Note that only one file can be moved at a time).

If the legend for a particular file is to be changed, simply doubleclick on the file to specify a new legend for that file.

The center part of the multiplot page is the plot control area, where placement of legend, frequency labels, plot ranges etc. can be controlled. The textfield at the bottom is for additional Gnuplot commands if additional labels, arrows or similar is to be included in the graph (refer the Gnuplot manual for information on gnuplot commands).

The resulting graph shown on the right part of the page can simply be copied or saved (right click on the graph) and if a postscript file is preferred (for use with Latex documents for instance) press the 'Save postscript image' button.

It is possible to save the data and plot definitions for the current work by pressing the 'save plot definition' button. The resulting file can then be reloaded at a later time for further work. It is important to note, that as Elchemea Analytical is a multi user system, it includes an automatic clean up facility which removes files (uploaded data files and generated image files) after one hour of inactivity. Thus if the user expects to take a break from the Elchemea Analytical system, remember to use the 'save plot definition' beforehand so as to not lose any work.

If the scaling of the data is not as intended, the scaling of the data can be changed by changing the 'z(x)' function definition (default is $z(x) = x$). To do so, simply write the new definition in the 'additional gnuplot commands' field. For instance if the sample has a 6 cm^2 area and the graph should report the impedance in area specific resistance, write the following in the text area:

```
z(x) = x * 6
set ylabel '-Z_{imag} ({/Symbol W} cm^2)'
set xlabel 'Z_{real} ({/Symbol W} cm^2)'
```

The first line changes the scaling and the two following lines change the axis descriptions to include cm^2 (leave them out if only the scaling are to be changed).

Chapter 5

File formats

Department of Energy Conversion and Storage at Technical University of Denmark
Impedance visualisation and analysis control software supports a number of different
file formats including gamry(R) and ZPlot(R) / ZView(tm) files.

The native file format is the i2b file format however. The file format is described below
with an example file:

```
Idun:/home/EFA/rig14/1255/4/s4_1007.i2b
Desc: 14test78_PS128052
:2009:01:17:23:17:30    1232230650    , Meas.:
```

```
37
82451 0.006685137356 0.00141940337
56173 0.0070404900979 0.0002431376363
38270 0.007484563409 -0.000686494133
26073 0.008028417169 -0.001394965951
17764 0.008634536255 -0.001947261525
12102 0.009370725592 -0.002396327231
8245 0.010187221398 -0.002722736196
5617 0.011033464434 -0.002911896725
3827 0.011876708228 -0.003031150569
2607 0.012699248889 -0.002996166924
1776 0.013426438402 -0.002909587586
1210 0.014135420185 -0.00274866887
824.5 0.014798469288 -0.002688357568
561.7 0.015415708783 -0.002545973023
382.7 0.016064610363 -0.002514285078
260.7 0.016722169513 -0.002355489951
177.6 0.017329395388 -0.002180886816
121 0.017877782005 -0.001969379374
82.45 0.018305518738 -0.001746990494
56.17 0.018698563455 -0.00151536534
38.27 0.01889247348 -0.001359174472
```

```
26.07 0.019155759436 -0.00132627701
17.76 0.019422716189 -0.001266016758
12.1 0.019661051995 -0.001295590415
8.245 0.019888830093 -0.001464146169
5.617 0.020308127403 -0.001725404958
3.827 0.020971000603 -0.001804639011
2.607 0.021602216143 -0.001638322228
1.776 0.02195975717 -0.00134448105
1.21 0.022350984057 -0.001060789257
0.8245 0.022530834712 -0.000697843738
0.5617 0.022641762282 -0.0004145079483
0.3827 0.0226543601828 -0.0003907922012
0.2607 0.0227450009778 -0.0003855313026
0.1776 0.0228211856604 -0.0001882806252
0.121 0.0229167510741 -0.0001616599266
0.0825 0.0229392097983 -0.0002138122947
```

The first 6 lines can contain text information (including meta information). Usually the third line contains a time stamp, but Department of Energy Conversion and Storage at Technical University of Denmark Impedance visualisation and analysis control software does not use this.

The 7'th line contains an integer describing the number of data points, and the following lines (nr 8 and onwards) is the actual data. Each data point is in the format:

Frequency Real_part Imaginary_part

Chapter 6

Installation and system maintenance

This chapter describes how to install or upgrade a Elchemea Analytical system.

6.1 Requirements

The Elchemea Analytical requires the following software to be installed on the target system prior to installation:

- A Linux type operating sytem (Only tested with CentOS-5®, but will likely work on other Linux type systems as well).
- Gnuplot® version 4.0 or later.
- The Apache® web server version 2.2.3-43 or later (it is possible that earlier versions of apache will also work, however this is not tested).
- The Perl® interpreter version 5.8.8 or later.
- The Perl Time::Hires module (installation of this vaires between distributions, thus needs to be installed manually, refer your deistrbution manual as to how).
- ImageMagick version 6.2.8 05/07/12 Q16 or later.
- GPL Ghostscript version 8.70 (2009-07-31) or later.
- Gnu 'make'. Other 'make' packages than the one from Gnu may also work, but has not been tested.
- A functioning connection to the internet. The reason for this is that Elchemea Analytical downloads and installs several Perl modules from CPAN.org durring the installation.

6.2 Installation

In order to install the Elchemea Analytical system, unpack the tar-ball in a suitable location, cd into the resulting ElchemeaAnalytical directory and run *make*.

Inspect the output of the make program and resolve any errors.

Once all errors have been resolved, run *make test* followed by *make install*.

In order to ensure that all servers start upon system reboot, add the following line to */etc/rc.local*:

```
/usr/local/bin/analytic/start_servers &
```

Finally, start up a web browser and point to this address

hostname.domain/cgi-bin/analytic/main.cgi (substitute hostname and domain with the appropriate values for your system) to check if the system is properly configured, the resulting page should look like figure 3.1;

6.2.1 SE-Linux

If SE-Linux is installed on the server and running in enforcing mode, the default configuration of SE-Linux will prevent the apache web server from accessing the LATEX server listening on port 4050 as well as the execution of various scripts in */usr/local/bin/analytic*.

To allow apache to connect to port 4050 as well as execute the scripts in */usr/local/bin/analytic/*, (on CentOS) execute the script *set_SELinux.rules.bash* found in the ElchemeAnalytics installation directory.

6.3 maintenance

Generally the Elchemea Analytical system requires little maintenance and the Elchemea Analytical system includes a facility for automatic software updates, to enable this, simply add the following line to root's crontab file:

```
0 8 * * 1 /usr/local/bin/analytic/analytic_updateer.pl >> /root/update.log.txt &
```

This will update the system once every monday. The automatic update system then fetches any new version which may have been deployed within the last week and installs this if it passes the software test (*make test*).

Chapter 7

Server structure

The programs mentioned in *italics* below all reside in the */usr/local/bin/analytic* and are written in Perl.

7.1 LATEX-server

The LaTeX server (LATEX-server) is responsible for compiling LaTeX reports of the fit results of individual fits. It honors the following commands:

- debug: Turns debug on and off.
- exit: Shuts down the server cleanly.
- unlink: Unlinks the specified filename. Note this is a potential security risk, so the LaTeX server should not be accessible from external sources!
- compile: compiles the document with the specified filename. The resulting pdf file is placed in *'/home/http/html/analytic/png/'*.

Chapter 8

System command interface (command line)

Although the Elchemea Analytical is designed to be used primarily through the web interface some programs can be accessed from the command line. Below is a list of the most used command line tools for the Elchemea Analytical system:

- */usr/local/bin/analytic/z_to_i2b*: This program converts Z-plot files to the 'i2b' file format.
- */usr/local/bin/analytic/gamry_to_i2b*: This program converts a Gamry® file to the 'i2b' format.
- */usr/local/bin/analytic/batchfit.pl*: This program can be used to fit multiple spectra to the same model. Call the program without any arguments to get a description of how to use it as well as a list of possible options.

Chapter 9

Module specifications

This chapter contains the module specification for the perl modules supplied as part of the ElchemeaAnalytic software suite. It includes function descriptions including number and type of any function arguments. Some of the modules are object oriented (with only a publicly accessible constructor) and in other cases the modules are function orientated.

In the case of function orientated modules, any functions exported by the module are described, both for what it does, as well as number and types of arguments.

In the case of the object oriented modules, any inheritance is also described (usually in the beginning of the module description). For the object instances, usually only the member functions intended to be public is described (as perl does not have a true private function declaration). Note that some of the object orientated modules define more than one class type, but as all the class types in this case behave similarly (polymorphic), only the main class is described as the subsequent class definitions implements the main class type behaviour.

Each module is described in it's own section.

9.1 Debug

Use: `my $id = Debug→new();`

This class is intended to be a base class for other classes to derive from so that easy debug functionality can be included.

Utility class for debugging. It contains the following member functions:

<code>\$id→debug()</code>	Sets or gets the debug level: level 0 is no debug, level 5 is complete debug including stack backtrace. This class only uses level 0 (no debug), level 1-4 (debug information displayed) and 5, debug info displayed with complete stack backtrace. The levels 1-4 lets other modules define debug levels inbetween the ones used here.
<code>\$id→writedebg(\$,[\$])</code>	Writes the string to standard error if debuglevel is 1 or higher. If override is specified (second argument which is optional), debug level 5 is assumed for this debug.
<code>\$id→die(\$)</code>	Appends stack backtrace to argument string and calls <code>CORE::die</code>
<code>\$id→print_setup()</code>	Prints out the complete current setup including all member functions and data fields (uses <code>Class::Inspector</code>).

9.2 SemaforeFile

Inherits from Debug (refer section 9.1).

This package makes file inout/output on multiprocess systems more easy by encapsulating file locking. To define a new semaforefile use the new method:

`my $id = SemaforeFile→new($filename);`

If the lockfile is not to be the default (`/var/lock/SemaforeFile/SemaforeFile.lock`) specify as second argument to new:

`my $id = SemaforeFile→new($filename,$lockfile);`

The package includes the following simple public methods on semafore files:

<code>\$id→readonly()</code>	Returns true if the file is readonly for the current user
<code>\$id→exist()</code>	Returns false if the file does not exists;
<code>\$id→filename()</code>	Returns the filename of the semafore file
<code>\$id→readlines()</code>	Returns the content of the file as an array with one line in each element Note that it removes any trailing newline from the read lines!

- `$id→writeline(@)` Writes the arguments to the file (NB: Overwrites file and add a newline to each argument if they do not already have it).
- `$id→append(@)` Appends the arguments to the file (Also adds newlines if nescesarry).

It is not nescesarry to check for file esistence in readlines as an empty array is returned if the file does not exist Note that the readlines function should only be used on small files as it globs the entire content to memory! For large files, use the more advanced member functions (see below). Also note that trailing newlines are removed from the individual lines. If this are not desired, use the `readline()` method described below.

The module also includes the following methods for advanced use: Note none of these functions check if the file exist before trying to open! The unsafe versions of open and close does not lock or unlock (assumes the user does this explicitly!)

<code>\$id→lock_ex()</code>	Locks file for exclusive use (Read, Write or Anppend)
<code>\$id→lock_sh()</code>	Locks file for shared access (Read only)
<code>\$id→lock_ex_nb()</code>	Locks file for exclusive use non blocking (Check return status!)
<code>\$id→lock_sh_nb()</code>	Locks file for shared access non blocking (Check return status!)
<code>\$id→unlock()</code>	Unlocks file
<code>\$id→open_read()</code>	Opens the file for reading (locks file shared if not already locked)
<code>\$id→open_readback()</code>	Opens the file for reading backwards (locks file shared if not already locked)
<code>\$id→open_write()</code>	Opens the file for writing (locks file exclusive if not already locked exclusive)
<code>\$id→open_append()</code>	Opens the file for appending (locks file exclusive if not already locked exclusive)
<code>\$id→close()</code>	Closes the file and unlocks it
<code>\$id→open_read_unsafe()</code>	
<code>\$id→open_readback_unsafe()</code>	
<code>\$id→open_write_unsafe()</code>	
<code>\$id→open_append_unsafe()</code>	
<code>\$id→close_unsafe()</code>	
<code>\$id→mtime()</code>	Returns the time of modification of the file as reported by <code>File::stat→mtime</code> , returns 0 if the file does not exist.
<code>\$id→readline()</code>	Reads and returns the next line from the file, assumes an open file Raises an exception (die) if not.
<code>\$id→fh()</code>	Returns the underlying file handle for direct IO (Use with care!)

Additionally the `$id→debug($)` member function (inherited from `Debug.pm`) can turn

debug information on and off \$id→debug(\$level) turns debug on and \$id→debug(0) turns debug off (\$level is the debug level, 1 - 5) This may be usefull if deadlock is encountered (so that the individual file locking operations can be monitored! If \$id→debug() is called without arguments it returns the status (i-e if debug in on 1 is returned else 0).

9.3 SocketClient

This module defines a number of communication functions used for accessing tcp:IP sockets on local and/or remote systems. The functions defined are listed below:

socket_client_raw(\$\$@)	Base function used by all subsequent functions, handles the raw tcp:IP cummunication. Arguemnts: server, port, [additional args to server]. The server can either be a ip-address or a hostname. Any additional arguments gets serialised with tab characters and 2 newlines are appended to the resulting string before transmission.
socket_client(\$\$@)	Same as above, but catches any communication errors in an eval guard.
socket_client_nocr(\$\$@)	Same as above, but do not append any newlines to the transmitted string.
socket_client_raw_nocr(\$\$@)	same as socket_client_raw() but do not append newlines.
serial_client(\$@)	communicates with a local serial server (which handles hardware communication on the serial port. Arguments: tty, args_to_server. The server is assumed to be the local server (either localhost or the public IP address of the server) and the port number is the tty number added to 202020 (Note wraparound!).
GPIB_client()	Communicates with the GPIB-server. Arguments are passed to the GPIB-server serialised with tab characters using socket_client_nocr(). The server is assumed to be the local server (either localhost or the public IP address of the server) and the port number is 12345.
serial_client_raw(\$@)	Same as serial_client() but without eval guard.
GPIB_client_raw()	Same as GPIB_client() but without eval guard.

9.4 Impedance::Header

This module specifies global file locations and other global variables used for the Elchemea-Analytical software package.

The module also specifies the colors of the resulting gnuplot figures (this is specified in the \$gpheader variable).

9.5 Impedance::IMPCGI

This module contains a number of utility functions for outputting properly formatted html code for user interface generation. Thus it mainly extends the CGI.pm module by Lincoln D. Stein. The module exports these functions in two groups.

The :html group exports these functions:

<code>print_header(\$[%])</code>	Prints the header information. Arguments: title. Any additional optional arguments (in the form of a hash) will be parsed along to the <code>header()</code> function supplied by CGI.pm. The function automatically appends a call to a javascript function logging users out after some time of no actions.
<code>print_end()</code>	Prints the help button and ends the html output with the proper tag.
<code>print_hidden()</code>	Prints a number of hidden fields used to maintain state.
<code>logout()</code>	Prints a logout button.
<code>action(\$)</code>	Prints a hidden field with an action parameter with the specified value which can be used for program control flow.
<code>EFA_start_html()</code>	A wrapper for <code>CGI::start_html</code> . Any arguments (in the form of a hash) are passed to <code>CGI::start_html</code> . Automatically appends a reference to the javascript source file on the server.
<code>js_back()</code>	Prints the javascript for going backwards (uses the <code>browser.back()</code> javascript call).
<code>get_CGI_value(\$)</code>	Retrieves the value of the specified CGI parameter (supplied by the web browser).
<code>get_CGI_value_clean(\$)</code>	Same as <code>get_CGI_value</code> , but does pattern match on the retrieved value and only returns the part that matches. The pattern match is <code>[\w\s\.\,]*</code> . This has the benefit of untainting the returned parameter value (For taint checks in perl and web access, refer Lincoln D. Steins book <i>Official Guide to Programming with CGI.pm</i>)

The :cgi group of functions include the following:

<code>get_CGI_value(\$)</code>	See above.
<code>get_CGI_value_clean(\$)</code>	See above.
<code>action(\$)</code>	See above
<code>menu_button(@)</code>	Prints a menu button. Arguments: name, value, style. The name will be the CGI parameter name, the value will be the text on the button and the style is a style class name to use for displaying.
<code>create_menu_field</code>	Prints the html tags to create a menu field.

<code>top_nav_bar_start()</code>	Prints the html tags to start the top navigation bar (table specifications etc.)
<code>top_no_button()</code>	Prints a no action button (goes nowhere) in the top navigation bar.
<code>top_nav_bar_button()</code>	prints a top navigation button. Arguments: File, name, value, style, [optional additional name, value and force triplets]. The file is the cgi-script to be called upon button press, the name,value and style arguments are passed to <code>menu_button()</code> and the additional optional arguments are used to initialise and print hidden html fields in the form of name-value pairs and a force argument (1 for force value, 0 for allow reuse of value).
<code>tab_newrow()</code>	Prints a new row in the top navigation bar.
<code>top_js_return()</code>	Prints a top navigation return button (uses the javascript printed by <code>js_back()</code> , see above)
<code>end_top_bar()</code>	Prints the end of the top navigation bar.

An additional function which is often used (but which is not exported by default) is the `format_model($)` function which parses a given impedance model and removes any empty serial or parallel connections as well as empty lines. The `format_model` function also removes any `"\r"` characters which may have been added by Microsoft based browsers. The function returns the resulting model.

9.6 Impedance::Base

Inherits from `Debug` (refer section 9.1).

This module defines all member functions which an `ElchemeaAnalytical Impedance` element must honor. Most of the functions are merely stups intended to be overloaded by the individual class definitions.

The module also includes class defininitions on the basic discrete elements (R, C, L and Q, Q beeing the constant phase element).

The module is intended to be used in conjunction with `gnuplot` but can be used as is (but no fitting will be possible).

To obtain a element instance call one of the constructors as show below:

```
$id = Impedance::Base→new($);
```

```
$id = Impedance::R→new($);
```

```
$id = Impedance::C→new($);
```

```
$id = Impedance::L→new($);
```

```
$id = Impedance::Q→new($);
```

All the constructors accepts a single argument which must be the impedance element number. It is advisable to make sure that element id's are unique in order to be able to

distinguish.

All Impedance element instances has the following member functions:

<code>\$id→type()</code>	Returns the type of impedance element.
<code>\$id→name()</code>	Returns the name of the element (usually the type and the id)
<code>\$id→nr()</code>	Returns the element id.
<code>\$id→function_name()</code>	Returns the Gnuplot function name of the element.
<code>\$id→value(\$[<i>\$</i>])</code>	Sets or gets the value of the specified element parameter (If 2 arguments are specified, the second is the value to be set).
<code>\$id→description(<i>\$</i>)</code>	Returns the description for the specified tag. valid tags are 'name' and the tags returned by the <code>tags()</code> member function. The 'name' tag returns a text string describing the whole element and the other strings describe the specified parameter. If called with no arguments, it returns the string for the 'name' tag.
<code>\$id→tags()</code>	Return a list of valid element parameter names.
<code>\$id→functions()</code>	Returns a string containing all the functions and variable declarations for gnuplot.
<code>\$id→F(<i>\$</i>)</code>	Returns the impedance value of the element at the specified frequency. The returned impedance is of type <code>Math::Complex</code> .
<code>\$id→fit()</code>	Returns a list of possible fit variables (all starting with value 1) for use with gnuplot.
<code>\$id→save()</code>	Returns a string containing functions to save the final fit variables (base value multiplied with fit variable). This is also for gnuplot use.
<code>\$id→print_line()</code>	Returns a string defining the impedance element. The format is described below.
<code>\$id→helperfunctions()</code>	Returns a list of parameter names for which special help functions exist for calculating useful estimates for start values for fitting. Note, that some impedance elements does not have any way of determining good start values in which case the <code>helperfunctions()</code> member function simply returns an empty list.

<code>\$id→helpfunction(@)</code>	This function has multiple uses. If called without any arguments, it returns the list from the <code>helper-functions()</code> member function. If called with one argument it returns a string describing how many additional arguments must be parsed to it (which may vary depending on impedance element) and in which order. If called with more than one argument, the first Argument is the parameter name to calculate start value for, and the additional arguments (which must be a string of the form: "\$frequency \$real_value \$imaginary_value") are used for the calculation.
<code>\$id→f_imax()</code>	Returns the frequency for which the imaginary value of the impedance is at it's maximum (negative) value. If the element type makes such a calculation invalid, the function returns undefined. If the frequency is either infinite or the DC case, the reported frequencies will be 1e100 Hz and 1e-100 Hz respectively. For frequency independent elements the function returns 0.

The `print_line` member function returns a string which can be used to reload an impedance element. The format is:

Type: `parameter1=value1, parameter2=value2,`

An exception to this is the complex elements (serial and parallel) elements which is defined and implemented in the `Impedance::Complex` class.

9.7 Impedance::RQ

Inherits from `Impedance::Base` (refer section 9.6).

This module implements `Impedance::Base` for R-C parallel connections as well as the R-Q parallel connection. (The Q beeing a constant phase element).

To obtain an instance, call one of the constructors:

```
$id = Impedance::RC→new($id);
```

```
$id = Impedance::RQ→new($id);
```

The module defines no additional member functions.

9.8 Impedance::W

Inherits from `Impedance::Base` (refer section 9.6).

This module implements `Impedance::Base` for diffusion type elements (W, O, G and T).

W is the Warburg element, O is the Finite length Warburg element, G is the Gerisher element and T is the 'Bounded Warburg' element.

To obtain an instance call one of the constructors:

```
$id = Impedance::W→new($nr);
```

```
$id = Impedance::O→new($nr);
```

```
$id = Impedance::G→new($nr);
```

```
$id = Impedance::T→new($nr);
```

The module defines no additional member functions.

9.9 Impedance::Complex

Inherits from Impedance::Base (refer section 9.6).

This module defines series and parallel connections of impedance elements. The resulting element can be treated as any other Impedance::Base derived element, thus it is possible using the serial and parallel elements to build arbitrary complex circuit layouts!

In effect the module defines two container types which from the outside behaves as a single impedance element.

To obtain an element instance, call one of the constructors:

```
$id = Impedance::Ser→new($id);
```

```
$id = Impedance::Par→new($id);
```

The module defines three new member functions:

<code>\$id→elements()</code>	Returns a list of Impedance::Base (or derived) elements.
<code>\$id→elements_all()</code>	Returns the complete element list for all elements in the container (recursively).
<code>\$id→element(\$)</code>	Returns the element instance with the specified id (usually obtained from the elements member function).
<code>\$id→add_element(\$\$)</code>	Adds an element to the list of elements contained within the element instance. The arguments are type and id of the element to add. It returns the element instance created. The element is created using the Impedance::Device::new_device(\$\$) factory function.

The `print_line()` function inherited from Impedance::Base is overloaded with slightly different behaviour. Instead of a string containing just a single line, the return value of `print_line()` contains multiple lines. The first line contains a start identifier (either '(' or '[' depending on type) and the last line contains the corresponding end identifier (') or ']'). All intermediate lines are obtained by calling `print_line()` on the individual element(s) in the container. Note that this may include additional complex elements resulting in

nested parantheses (Which is completely valid behaviour)!

Additionally the `tags()` function is also overloaded so that it returns a list fo the elements in the container instead of valid parameter names (as the complex element is a container it does not by itself have any parameters of which to set or get the value).

It is also not possible to define any help functions on complex elements, thus any calls to `helperfunctions()` merely results in the empty list.

Notice that `f_imax()` can only report summit frequencies between 1e-100 and 1E100 Hz. Thus if the true sumit frequency of the complex element used is outside this range it will be reported wrongly!

9.10 Impedance::Device

This module defines two functions, the first is `list_device_types()` which returns a list of valid impedance element type names.

The second function is a factory function: `new_device($$)`. It accepts two arguments, the first is an element ytpre name and the second argument is the element id (integer). The `new_device($$)` function returns the impedance instance created.

9.11 Impedance::Model

Inherits from `Debug` (refer section 9.1).

This module defines how to handle impedance models. It utilises the impedance elements defined by `Impedance::Base` and the derived classes.

To obtain an instance, call one of the constructors:

```
$id = Impedance::Model→new();
```

```
$id = Impedance::Model→new($data);
```

```
$id = Impedance::Model→new($model_ref);
```

```
$id = Impedance::Model→new($model_ref,$minf,$maxf);
```

The first constructor merely initialises a new empty model. The second constructor initalises an empty model but adds the impedance data in in the supplied data string to the model instance. The third constructor copies the model from the supplied model reference and copies the data from that instance as well. The fourth constructor copies as the third, but only copies those data which lies within the frequency range specified by the last two arguments! This can be used for partial fitting where only a specific frequency range is needed.

The individual impedance elements are stored in internal data structure which makes sure that only unique impedance element id's are used.

All `Impedance::Model` instance has the following public member functions:

<code>\$id→parse(\$)</code>	This function accepts a string defining an impedance model. The string must be lines of the format defined by the <code>Impedance::Base</code> function <code>print_line()</code> . The function parses this string and initialises the correct impedance elements based on this. The function returns the id number of the last impedance element added.
<code>\$id→print_model()</code>	Returns a string which can be parsed by <code>\$id→parse(\$)</code>
<code>\$id→device_types()</code>	Wrapper for <code>Impedance::Device::list_device_types()</code> .
<code>\$id→elements()</code>	Returns a list of impedance element names (nr).
<code>\$id→element(\$)</code>	Returns the impedance element specified. The function also gets elements from inside containers!
<code>\$id→elements.all()</code>	Returns the complete element list of the model. Recursively calls into any container elements. Note that unlike the <code>elements()</code> function, the actual device instances are returned as opposed to the element names.
<code>\$id→model.text()</code>	Returns a string representing the impedance model
<code>\$id→delete_element(\$)</code>	Deletes the specified impedance element.
<code>\$id→add_element(\$[\$])</code>	Adds an impedance element of the specified type. If an additional argument is specified, the id of the new element will be set to this number. If no second argument is specified, the next id is simply chosen. The function returns a reference to the added element.
<code>\$id→fit()</code>	Returns a list of possible fit parameters for current model.
<code>\$id→data([\$])</code>	Gets or sets the impedance data in the internal data field.
<code>\$id→delete_point(\$)</code>	Deletes the point specified from the internal data array. Returns the new data in the same way as <code>data()</code> . Note that index 0 is the point at the highest frequency.
<code>\$id→print_fit([\$])</code>	Returns a string containing all the functions necessary for gnuplot to initialise and fit the given model to the impedance data either in the specified filename or (in absence of an argument) in the internal data field. All data points are given even weight.
<code>\$id→print_fit_weight([\$])</code>	Same as <code>print_fit()</code> , however the actual fitting is done with uneven weight, so that data points with small values (absolute length of impedance vector) gets higher weight than points with larger values (weight inversely proportional to length).
<code>\$id→set_show_arcs(\$)</code>	Sets whether the plot functions should include the individual arcs in the plots or not. An argument of 0 disables the arcs, 1 includes them.

<code>\$id→print_plot([\$])</code>	Returns a string containing the function declarations and plot definitions for gnuplot to plot the specified data (if any) if no datafile name is specified, it uses the data in the internal data field.
<code>\$id→print_plot_range(\$\$[\$])</code>	Similar to <code>print_plot</code> , except it must have the min and max range specified as the first 2 arguments. The effect is that any labels are only printed if they have x-values within the specified range.
<code>\$id→print_bode([\$])</code>	Same as above, but for bode plots.
<code>\$id→print_plot_eps([\$])</code>	Same as <code>print_plot</code> , but for eps file output.
<code>\$id→print_bode_eps([\$])</code>	Same as above, but for bode plots.
<code>\$id→print_plot_error([\$])</code>	Returns a string containing the function declarations and plot definitions for gnuplot to plot the difference between the model and the specified data (if any) if no datafile name is specified, it uses the data in the internal data field.
<code>\$id→print_imp_sim(\$\$)</code>	Returns the function declarations necessary for gnuplot to plot an impedance plot of the current model. Arguments are the frequency range to plot (min and max).
<code>\$id→print_bode_sim(\$\$)</code>	Same as above, but for bode plot.
<code>\$id→F(\$)</code>	Returns the impedance of the current model for the specified frequency. The returned impedance is of type <code>Math::Complex</code> .
<code>\$id→subset(\$\$)</code>	Returns a subset of the data in the internal data field based on the specified minimum and maximum frequency.
<code>\$id→subtract_model()</code>	Returns the residual of the data in the data field once the impedance of the current model has been subtracted (subtraction done in the individual data points!).
<code>\$id→get_error()</code>	Returns the mean and maximum error for the data and model chosen. The error is calculated as the absolute difference of the data from the model both the real and for the imaginary part. The error is then normalised with the modulus of the data value. This is calculated for all frequencies in the data set and the mean and maximum values are returned (as percentages).
<code>\$id→minf()</code>	Returns the minimum frequency in the data set.
<code>\$id→maxf()</code>	Returns the maximum frequency in the data set.
<code>\$id→minr()</code>	Returns the minimum real part of the impedance in the data set.
<code>\$id→maxr()</code>	Returns the maximum real part of the impedance in the data set.

<code>\$id→mini()</code>	Returns the minimum imaginary part of the impedance in the data set.
<code>\$id→maxi()</code>	Returns the maximum imaginary part of the impedance in the data set.
<code>\$id→scale_factor([\$])</code>	Sets the scale factor function to the specified value. If no arguments, the scale factor function is set to the default 1, that is no scaling. Note that only simple proportionality scaling is possible.
<code>\$id→get_scale()</code>	Returns the current scale factor.
<code>\$id→set_limit(\$\$)</code>	Sets the plot limit of one of the following tags: 'xmin', 'xmax', 'ymin' and 'ymax' to the specified value. Arguments: tag, value.
<code>\$id→get_limit(\$)</code>	Returns the limit of the specified tag (see <code>set_limit()</code>).
<code>\$id→set_xlabel(\$)</code>	Explicitly sets the text string to be displayed in the xlabel.
<code>\$id→set_ylabel(\$)</code>	Explicitly sets the text string to be displayed in the ylabel.
<code>\$id→set_ylabel.bode(\$)</code>	Explicitly sets the text string to be displayed in the ylabel in bode plots.
<code>\$id→set_bode_mode(\$)</code>	Sets whether bode plot should only plot imaginary value or both real and imaginary (default). If called with a true argument mode is set to only imaginary values.

Additionally the following private member functions are also defined. Although Perl does not permit true private functions, do not use these functions from outside the class instances!

<code>\$id→print_plot_common()</code>	This function returns the function declarations for all elements in the impedance model.
<code>\$id→print_plot_main()</code>	This function does the actual work of <code>print_plot()</code> and <code>print_bode()</code>
<code>\$id→print_plot_main_range()</code>	This function does the actual work of <code>print_plot_range()</code> .
<code>\$id→print_plot_eps_common()</code>	Similar as above, but for eps output.
<code>\$id→print_simulation()</code>	This function does the actual work of <code>print_imp_sim()</code> and <code>print_bode_sim()</code> .
<code>\$id→save()</code>	This function saves the content of the data field to a temporary file (in /temp) It returns the filename of the temporary file.
<code>\$id→file()</code>	Returns the name of the data file.

Chapter 10

Web service interface

It is possible to use Elchemea Analytical as an web service. To do so call the following *ajax_model.cgi* web script with the parameters specified below. The script is located in *host.domain/cgi-bin/analytic/ajax_model.cgi* (substitute *host.domain* with the correct hostname and domain name of your Elchemea Analytical installation).

1. 'ajax', value: '1'
2. 'action', value: 'fit' or 'fitlist'. If fitlist is selected, the webservice returns a list of possible free fit parameters for the specified model
3. 'model', value: the impedance model as specified in chapter 3.
4. 'fitmode', value: 'Even' or 'Inverse'. Determines if data points have even weight (which is default) or weight is inversely proportional to absolute value of impedance (length of impedance vector). Optional parameter, only used for 'fit' action.
5. 'data', value: Impedance data in the form of multiple lines, each line in the form: frequency real_value imaginary_value
Note that the first line must be the highest frequency. Only used for 'fit' action
6. 'fitlist', value: comma separated list of free fitting parameters. The elements specified must be from the list returned by the fitlist action (refer item 2), only used for 'fit' action.

In the case 'fit' action is selected and the fit converges, the resulting response would start with the string 'OK' on a single line followed by the fit result. This would include arc summit frequencies as well as pseudo capacitances (in the case of RQ elements) as well as the full set of final parameter values. The output would also include references to a number of png files temporarily located on the web server (of approx 10 minutes or so). These images can be downloaded separately and shows the fit result and error plot for the data and model in question.

The file *webservice.html* found by pointing your browser to *host.domain/analytic/webservice.html* contains two web forms with the necessary form

elements to do a impedance fitting using the web interface as well as testing the 'fitlist' action. By submitting the 'fit' test form with the default parameters, the output should look like this (Your browser likely 'eats' the newlines, so use view source):

```

OK
L_1:1.509e-08
R_2:0.01823
R_3:0.009409
Y_3:0.0387
n_3:0.7735
INFO: Maximum frequency for arc for element RQ_3: 4442
INFO: Pseudo capacitance for arc for element RQ_3: 0.003808
R_4:0.00219
Y_4:3.358
n_4:0.8441
INFO: Maximum frequency for arc for element RQ_4: 53.64
INFO: Pseudo capacitance for arc for element RQ_4: 1.355
R_5:0.008561
Y_5:29.01
n_5:0.9397
INFO: Maximum frequency for arc for element RQ_5: 0.7008
INFO: Pseudo capacitance for arc for element RQ_5: 26.53
FITDATA

*****
Wed Dec 7 12:26:16 2011

FIT: data read from '/tmp/fitset_26083.dat' u 2:1:3:(1)
      #datapoints = 74
function used for fitting: h(x,y)
fitted parameters initialized with current variable values

Iteration 0
WSSR      : 1.32424e-05      delta(WSSR)/WSSR : 0
delta(WSSR) : 0              limit for stopping : 1e-05
lambda    : 0.00565434

initial set of free parameter values

fl_1      = 1
fr_2      = 1
fy_3      = 1
n_3       = 0.8003
fr_3      = 1
fy_4      = 1
n_4       = 0.7235
fr_4      = 1
fr_5      = 1
fy_5      = 1
n_5       = 0.9762

After 9 iterations the fit converged.
final sum of squares of residuals : 2.32063e-07
rel. change during last iteration : -8.43531e-07

degrees of freedom (ndf) : 63
rms of residuals (stdfit) = sqrt(WSSR/ndf) : 6.06922e-05
variance of residuals (reduced chisquare) = WSSR/ndf : 3.68354e-09

Final set of parameters      Asymptotic Standard Error
=====
fl_1      = 1.10797          +/- 0.008277      (0.7471%)
fr_2      = 0.97708          +/- 0.003211      (0.3286%)

```

fy_3	= 1.16728	+/- 0.1147	(9.828%)
n_3	= 0.773495	+/- 0.01046	(1.352%)
fr_3	= 1.01687	+/- 0.01714	(1.686%)
fy_4	= 0.519381	+/- 0.09599	(18.48%)
n_4	= 0.844051	+/- 0.04384	(5.194%)
fr_4	= 0.670101	+/- 0.05151	(7.687%)
fr_5	= 1.08185	+/- 0.01273	(1.177%)
fy_5	= 0.928554	+/- 0.01251	(1.347%)
n_5	= 0.9397	+/- 0.007901	(0.8408%)

correlation matrix of the fit parameters:

	fl_1	fr_2	fy_3	n_3	fr_3	fy_4	n_4	fr_4	fr_5	fy_5	n_5
fl_1	1.000										
fr_2	-0.623	1.000									
fy_3	0.485	-0.773	1.000								
n_3	-0.536	0.837	-0.992	1.000							
fr_3	0.510	-0.822	0.923	-0.930	1.000						
fy_4	-0.104	0.185	-0.319	0.305	-0.483	1.000					
n_4	0.225	-0.379	0.586	-0.566	0.742	-0.924	1.000				
fr_4	-0.317	0.520	-0.757	0.735	-0.870	0.767	-0.928	1.000			
fr_5	0.114	-0.196	0.314	-0.303	0.429	-0.755	0.720	-0.740	1.000		
fy_5	-0.102	0.177	-0.282	0.272	-0.380	0.633	-0.612	0.584	-0.441	1.000	
n_5	-0.075	0.128	-0.207	0.199	-0.288	0.555	-0.515	0.566	-0.876	0.140	1.000

END_TEXT
FILE:model_26083.png
REPORT:report_model_26083.png
ERROR:report_error_26083.png

Note that some of the numbers may be slightly different as a random number generator is involved. The last three lines indicates filenames, and in order to access the individual files, point your browser to *host.domain/analytic/png/* and select the file(s) specified.

Chapter 11

Troubleshooting

11.1 Server error is reported when starting Elchemea Analytical

- Inspect `/var/log/httpd/error_log`
- If SELinux is running in enforcing mode, try and disable it by using `'setenforce 0'`.
- If this has resolved the error, inspect the file `/var/log/audit/audit.log` and find the files/directories with conflicting SELinux labels. The likely culprits may be `'/var/lock'` and `'/var/semaphoreFile/'`, refer SELinux documentation as to how to do see and change labels.
- Reenable SELinux by running `'setenforce 1'` and check.

11.2 Model section of view is mangled

It is known that some versions of Microsoft Internet Explorer® display some of the HTML elements wrongly, thus leading to mangled model views. In this case either upgrade your browser, or switch to Firefox or Google Chrome (ElchemeaAnalytical has not been tested with Apple's Safari browser).

11.3 After fitting, pressing the report button only says 'no report ready'

- Check that the LATEX server is running. To do this type the following in a terminal:
`ps -ef | grep LATEX`
The output should look something like this:

```
sofc 10320 10297 0 10:17 pts/3 00:00:00 grep LATEX
sofc 19042      1 0 Aug02 ?      00:00:00 /usr/bin/perl /usr/local/bin/analytic/LATEX-server
```

If the last line is not observed, start it by calling
/usr/local/bin/analytic/start_servers as root.

- If the server is running, check that only one version of the LATEX.pm module is installed. Older versions for Risø Fuel Cells and Solid State Chemistry division fuel cell control system installed modules in a different location than Elchemea Analytic, and depending on search path, the Elchemea Analytic installer may not have discovered the older version and installed the new version independently. If this happens, the old version may be used by the LATEX-server, and unfortunately the old module misses some functions needed by the server.

To fix this, locate and delete the old module.

- If the problem persists (or only one version of LATEX.pm exists, then if SE-Linux is running in enforcing mode, then it may prevent the Apache webserver from accessing the LATEX-server on localhost port 4050. The easiest way to fix this is to run SE-Linux in permissive mode (nonenforcing). However, be aware that SE-Linux is part of the Linux intrusion detection system / security system, and thus running SE-Linux in nonenforcing mode may potentially expose the system to external threats it would otherwise be protected from. To run SE-Linux in non-enforcing mode in order to check if this is the cause of the problem, consult your Linux distribution manual (on CentOS version 6.x SE-Linux can be turned off by typing 'setenforce 0' in a root terminal).

The specific problem with SELinux is that Apache needs to be able to connect to port 4050 (the LATEX-server) and be able to read the corresponding PDF file (a separate issue). To allow apache to connect to port 4050 as well as execute the scripts in */usr/local/bin/analytic/* (on CentOS) execute the script *set_SELinux_rules.bash* found in the ElchameAnalytics installation directory.

11.4 Fitting does not finish (page displays 'working...' and stops)

Check that the 'utf8.def' file is included in your L^AT_EX installation, if not, locate the following line in LATEX.pm and uncomment it (place a # in front)

```
$res .= ' \usepackage[utf8]{inputenc}''\n' if ($utf8);
```

Note that this has to be done in the LATEX.pm file used by the implementation (Not the one in the install directory!).

11.5 Fitting takes too short time and no response is recieved

Likely your model contains one or more of the elements which contains a trigonometric function in the mathematical description (this includes Gerisher, Finite length Warburn and Bounded Warburg) and the fitting routine hit a value which resulted in infinite impedance (refer section 3.4).

11.6 Graphs not shown correctly and/or pages does not finish loading

Check that the default lock file (called SemaphoreFile.lock) for the SemaphoreFile.pm module has the right permissions. It is located in /tmp and should have permissions 666 (Yes, I know the number of evil...) During normal operation, it will be created with this permission, but sometimes the system may clean up the temp directory, and in this case sometimes it may be created with the wrong permissions. To resolve this, simply remove the file or manually set the right permissions (both operations may be necessary to do as root).

11.7 My screen is not wide enough to show all information

This can happen if you are using an older screen/projector which only allows a maximum horizontal resolution of 1024 pixels. To correct this, edit the file Header.pm in the Impedance directory (likely placed somewhere under /lib) and change the size variable from "set size 1.3,1.1" to "set size 0.9,0.7" as well as the pssize variable from "800x600" to "600x400" (Remember to change both variables!). Notice however that this is a site wide variable, so all users of Elchemea Analytic on this server will be affected.

11.8 Multiplot graphs are sideways

Make sure that gnuplot, ImageMagick and Ghostscript are up to date. On Centos/RHEL this can be achieved by executing the following commands as root:

```
yum update gnuplot
yum update ImageMagick
yum update ghostscript
```

Other distributions handle this in a different way, refer the distribution manual as to how to update software packages.

11.9 Multiplotting suddenly fails with an error message including the string 'all points y value undefined'

Due to the multiuser nature of the Elchemea Analytical system, uploaded files can not be saved indefinite on the server, and if a file has been left unused for some time (usually an hour) it will be deleted. This usually does not happen when working normally, but if the user leaves the Elchemea analytical session for extended time and then returns, this may happen (see section 4).

11.10 Some of the last tics on the graphs is missing (graph goes to 100 but tics only shown to 70 for instance).

This is caused by certain versions of Gnuplot. Version 4.2.x is known to do this. To correct this, edit the file Header.pm in the Impedance direstory (likely placed somewhere under /lib) and change the size variable from “set size 1.3,1.1” to “set size 1,1”. Notice however that this is a site wide variable, so all users of Elchemea Analytic on this server will be affected.