

# Plasimo User Guide

The Plasimo Team

April 23, 2014

# Contents

<b>1</b>	<b>Introduction</b>	<b>3</b>
<b>2</b>	<b>Downloading, Installing and Running Plasimo</b>	<b>4</b>
2.1	Downloading Plasimo . . . . .	4
2.2	Starting the Application . . . . .	5
2.3	Installing and running a model . . . . .	6
<b>3</b>	<b>The Graphical User Interface</b>	<b>9</b>
3.1	The Main Window . . . . .	9
3.2	The Global Settings Window . . . . .	9
3.3	The Model Editor Window . . . . .	10
3.3.1	The Tree Editor . . . . .	10
3.3.2	The Leaf Editors . . . . .	11
3.4	The Data Window . . . . .	13
3.4.1	Additional functionalities of the viewers . . . . .	14
3.4.2	Initial Viewers . . . . .	15
3.5	Model-specific GUI Extensions . . . . .	15
<b>4</b>	<b>Plasimo example models</b>	<b>17</b>
4.1	Introduction . . . . .	17
4.2	MD2D Model . . . . .	17
4.3	MD2D sputtering model . . . . .	21
4.4	MC model . . . . .	22
4.5	PLASIMO model - 1 . . . . .	23
4.6	PLASIMO model - 2 . . . . .	25
4.7	EM models . . . . .	26
4.7.1	EM model - 1 . . . . .	26
4.7.2	EM model - 2 . . . . .	26
<b>A</b>	<b>System requirements</b>	<b>29</b>
A.0.1	Requirements for Microsoft Platforms . . . . .	29
A.0.2	Requirements for GNU/Linux and Unix Platforms . . . . .	29
<b>B</b>	<b>Installation &amp; Configuration</b>	<b>32</b>
B.1	Downloading Plasimo . . . . .	32
B.1.1	CVS . . . . .	32
B.2	Building the GUM binaries . . . . .	34
B.2.1	Compilation on Linux/Unix Platforms . . . . .	34
B.2.2	Miscellaneous Compilers on Linux/Unix . . . . .	36

# Chapter 1

## Introduction

This is the Plasimo [1] user guide.

Plasimo is available for Unix and Windows users. For access please contact the Plasimo team [plasimo-devel@plasimo.phys.tue.nl](mailto:plasimo-devel@plasimo.phys.tue.nl). If Plasimo is suitable for your particular application, the Plasimo team will provide you a username and password for downloading the code. Before to give you access to download the code, we will be asked to sign a small contract for the software transfer.

## Chapter 2

# Downloading, Installing and Running Plasimo

Plasimo is available in two flavours: stable, released versions, and an experimental developer version, where all the changes are done for the next release. Releases are currently happening every couple of months and may be recognised by an even release number. So the development goes like version 2.4 (stable) → 2.5 (developer) → 2.6 (stable) etc. If fixes to stable releases need to be made, a fixed stable version may be released and an additional number will be used to indicate this fact; 2.6.0 is followed by 2.6.1 etc.

Before you get the Plasimo sources, you should take a decision which branch you want to use: “Stable” or “Developer” (also called HEAD). “Developer” is recommended for people who develop code for Plasimo or that need the latest features. The downside is that experimental changes are allowed to go in and carry the risk of breaking user input files, producing wrong results or crashing the user interface. Stable is normally more suitable for people that are mainly users of Plasimo, as care is taken to not break stuff for releases and users that need to update their input files are advised how to do so. The file `NEWS` contains the documentation on what has changed since the last release. Note that it’s possible to have more than one copy of the Plasimo source tree on your machine, so you can work with several versions if needed.

### 2.1 Downloading Plasimo

Assumes that you already have the rights for downloading Plasimo and you have decided which version of Plasimo you want to use. Depending on your operating system proceed to the relevant section below.

#### For Microsoft Windows users:

On the Plasimo’s web page <http://plasimo.phys.tue.nl> you can download the binary zip Plasimo using the username and password provided by the Plasimo team.

Go to ‘Manual and source code documentation’.

You can download the compressed zip file for Windows 2000/XP binary distribution of the chosen version of Plasimo. Download and extract the files on your hard drive. We recommend to extract the files in the main directory on your hard drive. A typical installation directory may be `C:\` (or `D:\`). The zip archive puts all files in a subdirectory with the name `Plasimo`, so you end up with a directory `C:\Plasimo`. This directory will be called the *Plasimo root directory* hereafter.

Note that you do not need to have administrator rights to install the modelling package, as long as you have write access to the installation directory. In particular, installing Plasimo does not require changes to the Windows Registry.

### For Unix/Linux users:

The following steps concern users who have CVS access to the code of Plasimo, as provided by Plasimo team.

Before to proceed with the installation and configuration of Plasimo, please be sure that you have checked carefully the system requirements and that you have installed all components mentioned in appendix A.

The following steps demonstrate how to check out and build an optimised version of Plasimo that includes graphical user-interface (GUI) support.

Firstly, open a shell (console) and change to the directory where you want to create the Plasimo subdirectory. Next type the following sequence of commands (replace **NAME** with your login name on the server). Please note that all commands and options under Unix are case-sensitive.

- `export CVS_RSH=ssh`
- `cvs -d :ext:NAME@plasimo.phys.tue.nl:/usr/local/src/cvs checkout gum`
- `cd gum`
- `./makeconf.sh`
- `mkdir linux-opt`
- `cd linux-opt`
- `../configure --prefix=~/.plasimo-opt`
- `make`

Assuming everything went fine, you are now ready to run Plasimo. In case any of the previous steps fails, please check if you have all prerequisites installed, or consult the appendix sections A and B for a possible solution.

Additional information and in-depth discussion of these download and installation processes are provided in appendix B. After building Plasimo, you may wish to fine-tune the behaviour of Plasimo, according to your personal preferences. This is also discussed in section B.

## 2.2 Starting the Application

### Execution on Windows:

In order to facilitate starting the executables under an MS Windows environment, a set of batch files is shipped with the distribution, these can be found in the Plasimo root directory.

You can start the version of Plasimo with the graphical user-interface by simply executing the script `Plasimo.bat`, which resides in the PLASIMO root directory, for example with Windows Explorer. Together with two console windows the PLASIMO main window appears. The second console window shows the log of PLASIMO. After starting the application with an 'empty' model, you should see a window like that in figure 2.1.

### Execution on Linux/Unix:

As part of the build process, some information about the location of various files is recorded and stored in two scripts, `set-plasimo-local` and `set-plasimo`. The first is used when one wishes to run the version of Plasimo at the location where it was build, the latter when one wishes to use an installed version of Plasimo.

Let us assume that we wish to run the version of Plasimo at the build location (`gum/linux-opt/`). This can be done by entering the following commands in the build directory:

- `. set-plasimo-local` or `source set-plasimo-local`

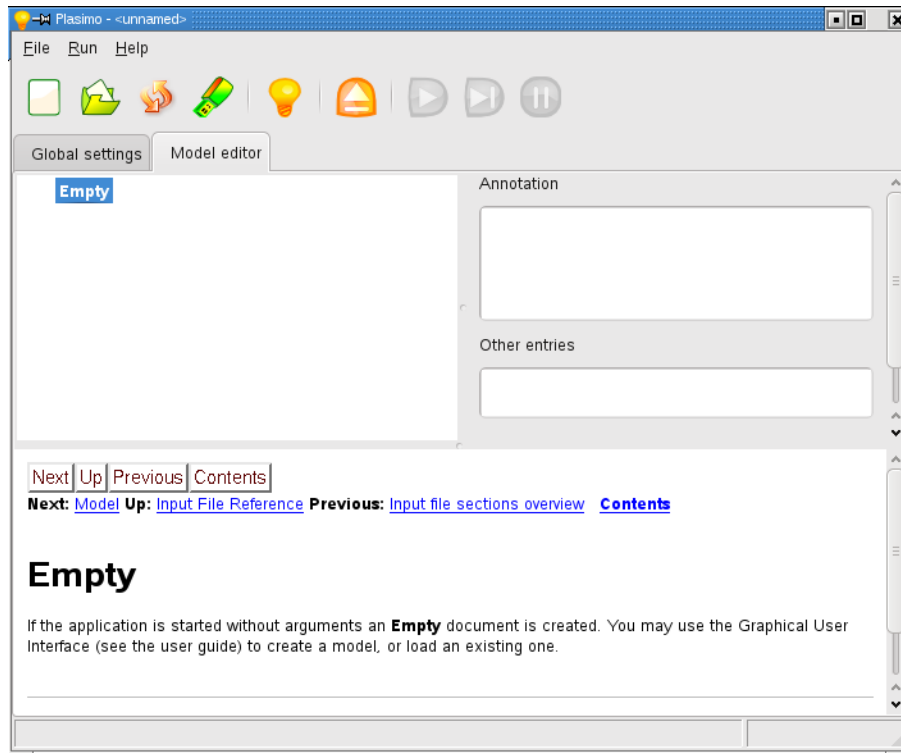


Figure 2.1: A screen-shot of Plasimo’s graphical user interface (GUI). The screen-shot has been taken just after starting the application, when no model file has been loaded yet. The elements of the interface are a menu bar, a tool button bar and set of tab windows. In the figure a tab window called ‘model editor’ is active.

- `./app/wx/wxplasimo`

After starting the application with an ‘empty’ model, you should see a window like that in figure 2.1.

## 2.3 Installing and running a model

In the application main window you see a menu. In order to allow quick access to the menu items, most commands are also available through the *toolbar*. Initially, the buttons **New**, **Open**, **Reload**, **Save**, **Help** and **Install** are active. While the icons are easily recognised, in case of doubt you may move the mouse over a button to see what it does: after a short time a ‘tooltip’ will be displayed. The remainder of the main window displays the properties of the active model. We will get back to that later.

1. **New** button gives you the possibility to create a new model by selecting a model type from the pop-up menu. The procedure how to create a completely new model is explained step by step in chapter 4.
2. **Open** pops up a file dialog which allows you to load a model by selecting an existing input file. A number of existing models are shipped with the Plasimo distribution. All the Plasimo input files reside in the directory `Plasimo/input/`. The files in this directory are separated in sub-folders depending on the sub-modules of Plasimo. Some of the existing example input files are described in more detail in the chapter 4.

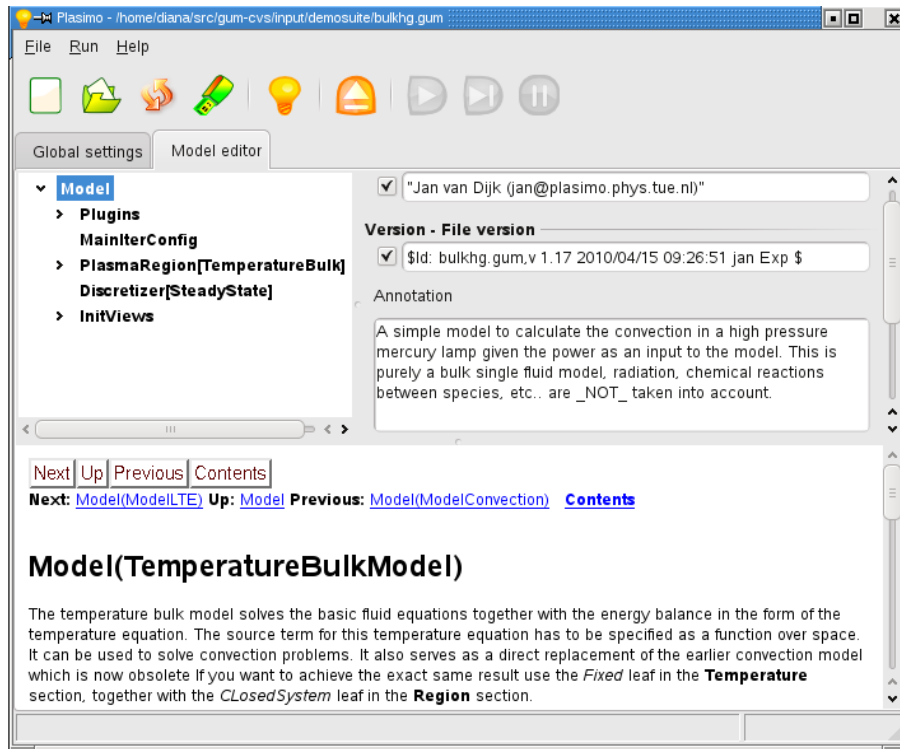


Figure 2.2: A screen-shot of Plasimo’s graphical user interface (GUI). The screen-shot has been taken just after loading an existing model, named `bulkhg.gum`.

3. Please load the example file `Plasimo/input/demosuite/bulkhg.gum`. If there were no errors, the main window should now look like 2.2. In the tree window on the left you see a tree representation of the `bulkhg.gum` model. The window on the right shows the contents of the active tree item. These editors will be discussed in detail in chapter 3, tutorial sessions are provided in chapter 4.
4. The **Reload** button allows you to reload an input file. Plasimo uses simply text files that can be edited with any of your favourite editors. In case you made changes in the this text input file, this provides a quick way to reload the modified model.
5. The **Help** button activates or deactivates the visibility of a help window on the bottom part of the screen. By default it is activated. It shows you a detailed description and requirements of the active tree item.
6. After a model has been loaded or modified to the user’s needs, it must be prepared for running. Press the **Install** button to do this. This will do all kinds of necessary one-time installations, and perform various checks on the sanity of the model file. Installing a model might therefore take some time, typically up to a few seconds. If anything goes wrong during installation, a message box will be popped up, telling the user what went wrong. After the problems have been solved, one can try to install the model once more.
7. When the model is successfully installed, additional buttons are activated. These are:
  - **Run**. Run the successfully installed model.
  - **Proceed (one step)**. All models are assumed to run in steps. A step can be either a time step, or an iteration. Selecting this item causes a (paused) model to do one such step. This can be useful if you use the application interactively, for example to find out the reasons for divergence of otherwise unexpected behaviour.

- **Stop.** Pauses a running model; this can be useful to inspect intermediate results.

For more information how to modify an existing model and how to manipulate the input data, please follow the steps in 4. Information about all the functionalities of the graphical user interface of Plasimo, please read chapter 3.



## Chapter 3

# The Graphical User Interface

This chapter discusses the usage of the graphical user interface (GUI), its functionalities such as the various model editors, settings and model specific GUI extensions.

### 3.1 The Main Window

After starting the application you should see a window like that in figure 2.1. This *application main window* contains a number of elements, from top to bottom we have:

- A menu bar, containing the menus **File**, **Run**, **Help**
- A tool bar, discussed in section 2.3
- A set of *tab windows*, named **Global settings** and **Model editor**
- A (HTML) **Help** window, on the bottom part of your screen, showing a detailed description and requirements of the active tree item.
- A status bar, displaying the current status of the model **Installed**, **Paused**, **Running**, etc.

As the menu/tool bars were previously discussed in chapter 2.3, the following text provides detailed information about the tab windows **Global settings**, **Model editor** and the additional **Model Data** window, that appears after installing a model, as well as some model specific extensions.

### 3.2 The Global Settings Window

In the graphical user interface the contents of the global configuration file are visualised in the tab window **Global Settings**. It is important to notice that some of the settings in the configuration file take effect only after the application is closed and restarted, while others take effect while it is still running. Among other things this can be used to fine-tune the behaviour of Plasimo (changing the appearance of plot data and the like).

In the interface the active section is explained in the HTML help window. For an overview of all available sections, we refer to the Plasimo user reference guide.

Most people can use plasimo without ever needing to use this window. Therefore, we shall continue this text with a discussion of the model editors and how you can edit a model using the GUI.

### 3.3 The Model Editor Window

Initially, the **Model editor** window has been activated. It is one of the tab windows previously mentioned and allows the user to create new model or edit existing one. Only one Plasimo model can be active, multiple documents are not supported.

In general, the structure of the **Model editor** window resembles the interface of most file managers, showing the directory (section) tree on the left (**Tree editor**), while the window on the right shows the contents of the active section (**Leaf editor**).

#### 3.3.1 The Tree Editor

The tree view shows the *nodes* or *sections* which are at the basis of the hierarchic structure of Plasimo input files. Please note that the structure is *recursive*, that is, sections may contain (sub)sections themselves. Each section contains a group of conceptually related subsections and variables.

A section can be activated by clicking the left mouse button on the section entry in the tree. The variables which belong to the active subsection are shown in the ‘section window’ at the right hand side. Furthermore, in the **Help** window at the bottom some information about this section is displayed.

If you want to add a subsection, click the right mouse button on the section to which you want to add it. This will pop-up a context sensitive menu which allows you to make an appropriate selection. This way it is guaranteed that you add subsections only at locations where the model will actually look for them. By clicking on the check box with the RMB, a pop-up menu appears which offers the following choices:

- **Cut** Removes the active node and the contents is copy to the clipboard.
- **Copy** Copy the active node
- **Paste** Paste the copied/cut node
- **Paste into section** Paste a copied node into the active section
- **Disable** this disables the entry if it is enabled. Disabled sections can be recognised by the light-up colour
- **Enable** this enables the entry if it is disabled. Pressing the right mouse button over a disabled section pops up a menu which allows to re-enable the section.
- **Undo all changes** This option will undo the results of editing the active node. Node that after switching to another node, editing can no longer be undone.
- **Add...** Add a relevant context dependent section. It is possible to have multiple choices. After one is selected a small addition window appears with the following options:
  - **Create default section** This will create a section with the default specifications. Note, that the relevant subsections will not be created recursively.
  - **Copy file into a section** This option allows you to copy data structures from external files. It works like this: the model will copy the data from an external file and paste it in your input model file.
  - **Include file as a section** This option also allows the usage of external data files. But it works slightly different: the model will only use the external data without writing the whole data it in your model file. Instead, you will see in your input file in the relevant section a line like this: `Include filename.in SectionName`. This allows to have only one definition of an argon mixture, say, on disk, which can be shared (included) by all models that describe some argon plasma.

### 3.3.2 The Leaf Editors

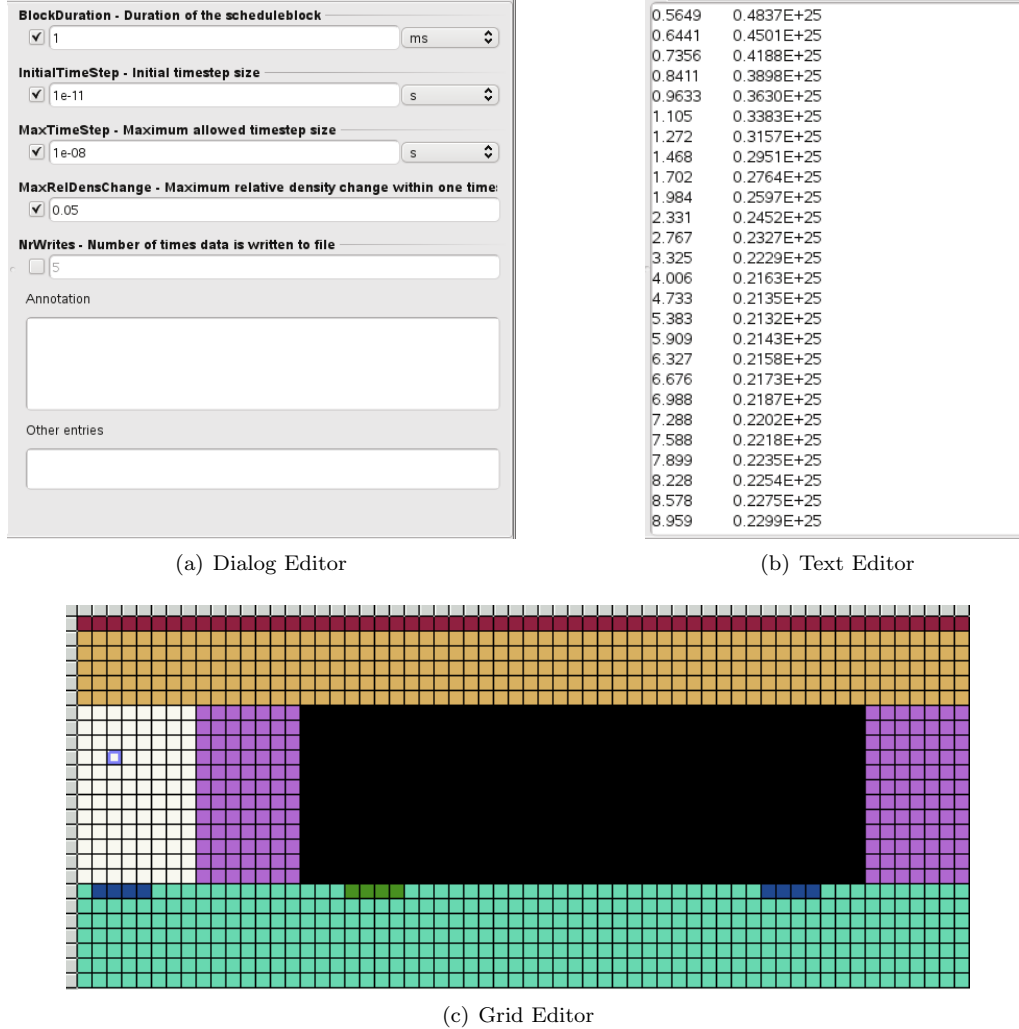


Figure 3.1: The various leaf editor windows. Figure 3.1(a) shows the dialog editor. The example shows the time settings of a time-dependent model edited using the dialog editor. Figure 3.1(b) shows a text editor, as an example is a lookup table. Figure 3.1(c) shows the grid editor in action (here a PDP geometry is being edited).

At the right-hand side of the model editor window, an appropriate editor for the data items of the active node are shown. Depending on the node contents, you will see a **Dialog editor**, a **Grid editor** or a **Text editor** window. In figure 2.2 a dialog editor is visible. The various editors are displayed in figure 3.1 and will be discussed next.

#### The Dialog Editor

The node dialog editor window (figure 3.1(a)) is the one you will most commonly encounter. It contains one *leaf block* for each data item type. The name of the data item and a short description are displayed above the box. Note that a box may be empty. Clicking on the name of the data item with the right mouse button (RMB) pops up a menu which allows you to add a new leaf of the relevant type. The default values will be used.

Each entry has a check box on the left, followed by one or more editor or choice fields. By clicking on the check box with the left mouse button (LMB), the entry can be disabled or enabled. Disabled entries are not taken into account when the model description is interpreted by Plasimo. They appear in grey and cannot be edited.

In addition, two small text editor windows are available. The 'Annotation' widget can be used to provide a description of the active section. If one exists, it will be shown. The window titled 'Other entries' can be used to add entries in text raw form. Plasimo also dumps entries it cannot handle here when creating a dialog editor, so it should be empty after creating a fresh dialog editor.

## The Grid Editor

The grid editor (figure 3.1(c)) facilitates editing of grids ("matrices") consisting of elements which can be represented as strings. Sometimes the matrix is too big to be visualised in the grid editor. You can zoom out/zoom in the matrix by pressing the control key and scrolling the mouse wheel. The matrix consists of cells with different colours as to each colour is assigned different material type.

There are two editing modes, the *Grid layout mode* and *Grid cell mode* mode. By default, the cell mode is active. One can switch between these modes via the context menu which is activated by pressing the right mouse button on the grid region.

The *Grid cell mode* deals with the properties of the cells, such as assignment of new material type. Clicking or selecting cells with the left mouse button (LMB) will assign the active colour to them (by default this is the colour black, which always corresponds to the material with the lowest alphabetic key value). If the control key is down, instead the material under the LMB is picked up as active colour. New material colour-index pairs can be created via the menu which is activated by clicking the right mouse button and selecting **Select material**. A list of previously defined materials will pop up for selection of material type.

In *Grid layout mode* only entire rows or columns can be selected. Row and column selection can be activated by clicking on the grey label bar at the left and top side of the grid, respectively. The presently selected row(s) or column(s) can be cut and copied via the menu which is activated by clicking the right mouse button. Then, the cut or copied row(s)/column(s) can be pasted before the active row/column.

Finally, after editing the geometry matrix, you may want to keep or discard the changes. The changes will be automatically saved after switching to another node. If you do not want to save the changes you made, you can retrieve the original matrix by pressing the RMB on the **GeometryMatrix** in the tree editor and selecting **Undo all changes** from the menu. Note, that after switching to another node, editing can no longer be undone.

## The Text Editor

The text editor window allows the user to manipulate the data entries in raw format. In some cases the creation of a text editor window is requested explicitly, because the nature of the data does not allow convenient editing inside a node dialog or grid editor dialog. Perhaps an appropriate editor will be provided later. An example is a look up table in a two-column format (see figure 3.1(b)).

The text editor is also used as fallback option if the creation of a node dialog or grid editor window fails for some reason. If the latter happens, the user will be informed and is kindly requested to send a bug report to the Plasimo maintainers.

There are a few things to think of when using the text editor window. Firstly, disabled leaves occur in raw text format with an underscore (-) prepended. Secondly, make sure to quote items inside this entry which consist of multiple words. As an example, an entry which specifies a name should be written as **Name "Jan van Dijk"** instead of **Name Jan van Dijk**. In the latter case, you will perhaps get an error when Plasimo interprets this entry, but more likely the part **van**

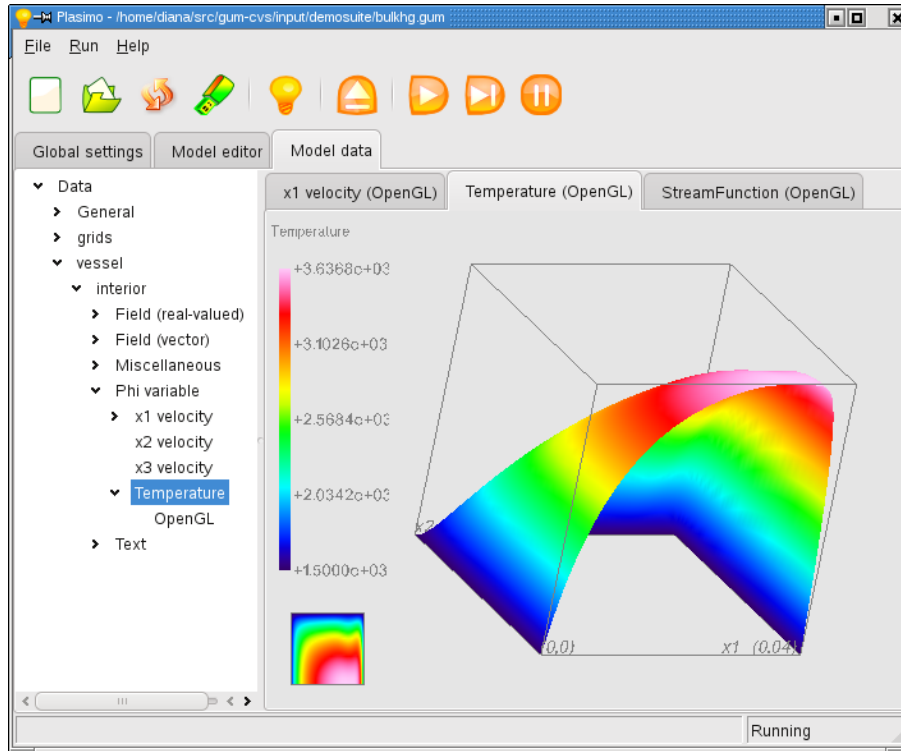


Figure 3.2: A screenshot of Plasimo’s graphical user interface (GUI). The screenshot has been made during a run of the model `bulkhg.gum` and shows an OpenGL plot of the temperature of the plasma region.

Dijk will be silently ignored. The text control correctly handles special characters, so newlines, tabs and backslashes can be safely used.

### 3.4 The Data Window

After installing the model, an additional tab windows has appeared, named **Model data**. You will see that the available data in this window have been organised as a tree, displayed on the left. The tree consists of a model-dependent data, which may be grouped into sections of related variables. As a minimum, a section called **General** has been created for every Plasimo model. Not surprisingly, its contents provide some general information about the model, such as the status of the calculation (you may think of data like elapsed time and the number of iterations). This can be activated by double clicking on the item **Common data** inside section **General**.

For most of the Plasimo models, of particular interest is the section **Interior - Phi variable**, which contains some important primary plasma parameters like the bulk velocity, pressure, densities.

For each data item a “viewer” can be created. Creating a viewer can be easily achieved by moving the mouse over the data item ‘Temperature’, say, and double click on it will open the default OpenGL viewer and you get the result shown in figure 3.2. The viewers are visible at the right hand side of the data window, and are displayed as a series of tab windows. Please take a moment to see what data are there.

Additional viewers are available and you can choose one by clicking the right mouse button in the selected item. This will pop up a menu which offers the option ‘Add view’. Some additional functionalities of the various viewers are discussed in the following section.

### 3.4.1 Additional functionalities of the viewers

Some additional functionalities of the various viewers are available. The next text discusses the different viewers available in Plasimo. Depending on the data item different viewers are available. Plasimo will offer you the appropriate suggestions for the relevant data item.

#### OpenGL

OpenGL viewer is the default viewer in Plasimo for plotting 2D and 3D plots. Using the left mouse button and dragging on the plot you can rotate it in all the possible directions. The plot window can also be controlled with the following keys:

- f – Switch between solid and wire frame plot.
- r – Reset the configuration parameters to their default values.
- l – Toggle lighting.
- s – Toggle smoothing.
- o – Include the origin in the plot.
- p – Print the plot window to a postscript file.
- d – switch between the directions
- c – switch between the components
- C – Toggle enabling (all) clipping planes
- <, > – Decrease/increase the number of planes that is drawn (3D only)
- X,Y,Z – This is available only for 3D plots. Draw planes with constant first, second or third coordinate.
- x , y , z – Activate the x, y or z coordinate. All subsequent coordinate-specific commands will affect that coordinate. The commands below are all coordinate-specific:
- D — Open a dialog window that allows you to configure the way the active coordinate (x, y or z) is handled inside the plot. You can control the coordinate range and the usage of a log scale. A help window is available inside the dialog window.
- a – Increase the position of the lower clipping plane.
- A – Decrease the position of the lower clipping plane.
- b – Increase the position of the upper clipping plane.
- B – Decrease the position of the upper clipping plane.

#### MathPlot WireFrame

This viewer is suitable for plotting data in a X-Y format. This is the default viewer in Plasimo for plotting this type of data. Pressing the right mouse button, the pop up menu will give you a list of all options available for this plot.

#### UC WireFrame

This viewer is an other option for plotting data in a X-Y format. Pressing the right mouse will give you the useful option to export the data of the plot as ASCII.

#### PlotMtv

Choosing this option, Plasimo will prepare the data to be plot using PlotMtv program. Of course, this requires that you have installed the program PlotMtv. Then the plot will be visualised in external PlotMtv window. The viewer is suitable especially for vector plots.

#### Grid

This viewer shows the data for one field variable as a spreadsheet. The cells in the spreadsheet are placed to resemble the positions of the corresponding data field variable in the master grid. This means that for a cell showing the value of a field defined at the nodal points (e.g. the species

densities), the north, south, west and east cells are empty, because in the mastergrid they represent the positions of the flux field variables. See the grid document for more information.

### 3.4.2 Initial Viewers

Sometimes, in the working process you may need to install the program multiple times and to monitor some data item at different conditions for example. After each Install of the program you have to open the relevant viewer to monitor the data item in run time. This could be annoying sometimes, so that Plasimo offers the option to avoid this by setting initially the viewers of the variables you want to monitor. As a result, when you install the model, the viewers will be initially open with the set of specifications.

Assume that you want a viewer for "Temperature" data item, as it is shown in figure 3.2, to be initially open always when you install the model. This can be done by adding the optional section **InitViews** in the model tree. Press with right mouse button on the **Model** and select **Add InitViews** from the pop up menu. Next, you have to add the relevant subsection: press the rmb on **InitViews** and add default section **ViewList**. On the leaf dialog editor on the right hand side of the window press again the rmb on the Viewer leaf to add one. Adding a new viewer leaf will displays 4 empty boxes. In the first one you have to specify the data location, for example *vessel/interior/Phi variable* In the second box you have to specify the name of the data item *Temperature*. In the third you can specify the viewer type. If it is empty, the default viewer will be open. In the last box optionally you can specify some attributes, such as min and max values of the x, y or z coordinate and logarithmic scale. All the attributes must be separated by comma, for instance: `xmin=0, xmax=5, ymin=3, ymax=20, zmin=1e12, zmax=1e20`, You can add viewers leafs as many as you can/want. Note that the editor is case sensitive!

## 3.5 Model-specific GUI Extensions

In section 2.3 it was mentioned that in addition to the functionality which is shared by all GUM simulation types, a number of application-specific extensions may be present. There are two such extensions available:

- Grid resizing: the grid can be re-dimensioned on the fly;
- Adjustment of  $\Phi$ -variables: Plasimo models allow the inspection and adjustment of the iterative control parameters of the convection-diffusion ( $\Phi$ ) variables while a simulation is in progress.

If the model supports these functionalities, after installing it, you will see an additional file menu called **Extra**. This menu manages the specified functionalities.

### Grid Resizing

Grid resizing can be seen as a *poor man's* implementation of the *multi-grid algorithm*. Essentially, in such methods the number of grid cells is not kept constant. Rather, the results of rapidly converging but inaccurate calculations on coarse meshes are combined with accurate, yet slowly converging calculations on finer meshes. The result is an net reduction of the time to convergence. Typically a number of grid changes is made in both directions. A full multi-grid algorithm has not been implemented in Plasimo. However, wxPlasimo allows its users to manually adjust the mesh size while a simulation is in progress. After a grid resize all grid-dependent variables are interpolated to the new grid and the iterative process resumes.

Please note that this grid resizing can also be configured to be done automatically at the input file level, see section MainIterConfig of chapter 'Input File Reference' of the User Reference Guide for details.

### $\Phi$ -Variable Adjustment

This allows the adjustment of the iterative control variables of the  $\Phi$ -variables which are part of the model. More specifically, the user is allowed to set the actual value of the  $\Phi$ -variables' *under-relaxation factor* (urf). The urf is the fractional amount of the field correction which is determined in an iteration which is actually applied to the old value of the field. Hence, when urf is unity the correction will be completely applied, while for smaller values only a partial adjustment is done. There is a trade-off between speed and robustness here: while smaller under-relaxation factors tend to stabilise the simulation, the time-to-convergence may be considerably longer.

If you select "Phi-Variable Adjustment" from the **Extra** menu in run time, a dialog appears which allows you to select a  $\Phi$ -variable, if any are present. If one is selected, a second box appears which allows to change the value of the under-relaxation factor, the actual value is proposed as default. Note that in Plasimo, depending on the setting for the urf-adjustment, the relaxation factors are continuously adjusted. Also a urf you set in this dialog is subject to later change.

Secondly, the option 'Freeze' can be activated. The effect is the same as that of a zero under-relaxation factor: no update is done whatsoever. Prefer to enable Freeze above setting a zero urf: in the former no calculation will be done for the variable, considerably saving computing time. Additionally, the discretisation of some of the variables relies on the assumption of a non-zero urf.

Of course freezing the variables cannot lead to physically realistic results, this option should therefore not be used in production work. Rather, it is meant to facilitate the analysis of the physical inter-dependencies between the model equations; in particular it can be useful in the study of convergence problems in the model development stage.



# Chapter 4

## Plasimo example models

### 4.1 Introduction

This chapter explains some of the existing demo models that were shipped with the Plasimo distribution. The examples give also information how you can edit an existing model and create a new one using the graphical user interface.

### 4.2 MD2D Model

In this section an example input file is used to show how the MD2D module is used.

**Example input file:** `demo.md2d`

**Location:** `input/md2d/`

This input file contains a model of a low pressure DC discharge with a simple geometry: two parallel electrodes at a distance of 50 cm, enclosed in a glass tube with a diameter of 5 cm. To keep the model as simple as possible we will start using Cartesian coordinates. The gas is He and only electron impact ionization is considered. The gas pressure is constant at 1 Torr, and a constant voltage of  $-400$  V is applied. You can find the input file `demo.md2d` in directory `input/md2d/`.

After installing the model, an additional tab “Model Data” appears in the main window. On the left hand side the in the herarchial tree the following sections are present:

- **Discharge Region:**

This section contains all the calculated variables; the section is subdivided into:

- Electron energy - this subsection contains the data from the energy equation, such as the mean electron energy, mean energy source, mean energy flux density.
- Reactions - contains the rates of the specified reactions.
- Species - contains the calculated variables for each species, such as density, mobility, source, flux.

An OpenGL plot of a variable can be activated by double-clicking a variable in the tree panel on the left.

- **EM Region:**

In this section you can see the EM data, such as potential distribution, electric field, surface and volume charge densities.

An OpenGL plot of a variable can be activated by double-clicking a variable in the tree panel on the left.

- **General:**

Naturally contains the general, common data, such as the status of the simulation. You can activate the **Model Status** by double-clicking on **Common data**.

- **Geometry:**

Here you can see the geometry - the configuration and the control volume grids.

## Output files

All calculated variables are not only shown in the GUI but are also stored on disk. The data path as well as the number of writes of the output data can be specified in the input. (see **Schedule**) By default, the program store the output files in the main plasimo folder if no data path is specified. The output files:

**history.out:** gives the calculated variables as a function of time. The averaged quantities are written with a frequency specified by the user.

**info.out:** the averaged values, written only once at the end of the simulation.

**info.txt:** the same data as in **info.out**, but written with the user-specified frequency.

**reaction\_analysis.out** and **reaction\_analysis.txt:** detailed reaction analysis: production and destruction contribution of each reaction for each species. The files are written like **info.out** and **info.txt**.

The following files represent the spacial distribution of the quantities. The data are written as many times as specified in the input file.

<b>n00.txt</b>	electron energy density [ $\text{J m}^{-3}$ ]
<b>phi00.txt</b>	electron energy flux density [ $\text{W m}^{-2}$ ]
<b>S00.txt</b>	electron energy source [ $\text{W m}^{-3}$ ]
<b>D00.txt</b>	electron energy diffusion coefficient [ $\text{W m}^2$ ]
<b>mu00.txt</b>	electron energy mobility coefficient [ $\text{J m}^2 \text{V}^{-1} \text{s}^{-1}$ ]
<b>Relas00.txt</b>	rate of electron energy loss from elastic collisions [ $\text{W m}^{-3}$ ]
<b>epsilon.txt</b>	mean electron energy [J]
<b>n01.txt</b>	density for species 1 [ $\text{m}^{-3}$ ]
<b>S01.txt</b>	source for species 1 [ $\text{m}^{-3} \text{s}^{-1}$ ]
<b>D01.txt</b>	diffusion for species 1 [ $\text{m}^2 \text{s}^{-1}$ ]
<b>mu01.txt</b>	mobility for species 1 [ $\text{m}^2 \text{V}^{-1} \text{s}^{-1}$ ]
<b>phi01.txt</b>	flux for species 1 [ $\text{m}^{-2} \text{s}^{-1}$ ]
<b>R00.txt</b>	reaction rate for reaction 1 [ $\text{m}^{-3} \text{s}^{-1}$ ]
<b>K00.txt</b>	reaction rate coefficient for reaction 1 [ $\text{m}^3 \text{s}^{-1}$ ]
<b>Pp01.txt</b>	power dissipation for species 1 [ $\text{W m}^{-3}$ ]
<b>P.txt</b>	dissipated power density [ $\text{W m}^{-3}$ ]
<b>J.txt</b>	current density [ $\text{C s}^{-1} \text{m}^{-3}$ ]
<b>V.txt</b>	potential [V]
<b>E.txt</b>	electric field [ $\text{V m}^{-1}$ ]
<b>Er.txt</b>	reduced electric field $E/p$ [ $\text{V m}^{-1} \text{Pa}^{-1}$ ]
<b>E_N.txt</b>	reduced electric field $E/N$ [ $\text{V m}^2$ ]
<b>rho.txt</b>	volume charge density [ $\text{C m}^{-3}$ ]
<b>sigma.txt</b>	surface charge density [ $\text{C m}^{-2}$ ]

The numbering of the species is the same as the order of the input file. i.e. in this case **n01.txt** contains the density distribution of the electrons and **n02.txt** the density distribution of the second defined particle, in this case  $\text{He}^+$ . The same numbering applies for the other variables.

## Results and discussion

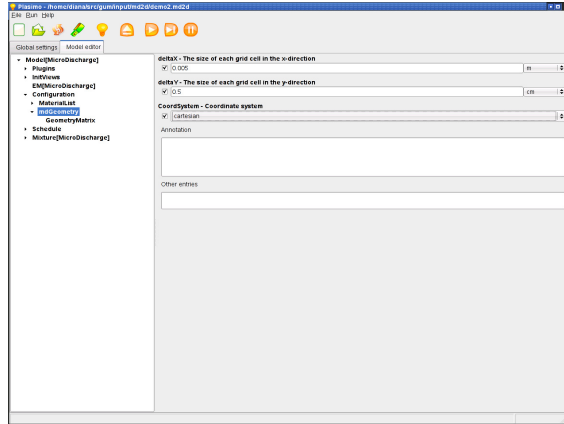
When the model is loaded, installed, and run, it will be possible to identify several regions in the discharge:

- Cathode fall region (Cathode dark space):  
Most of the potential drop between the electrodes occurs in the cathode dark space. This is a region with a strong electric field due to the positive space charge in front of the cathode. The strong field accelerates ions toward the cathode and electrons toward the negative glow region. The electrons ejected from the cathode are accelerated and gain energies up to the cathode fall.  
Select the **EM region** node for the **Potential**, **Ex**, and **Volume charge Density**, and the **Discharge Region** for the **Flux x** of the ions and electrons, and the mean electron energy.
- Negative glow:  
In the negative glow an intense electron impact ionization occurs. The electrons lose their energy and the E-field drops to almost zero. Select the **Discharge Region** node for the reaction rate and the power dissipation of the electrons.
- Faraday dark space:  
As the electrons have dissipated their energy, excitation and ionization will become less and less frequent, because electrons do not gain energy in the weak field. In this region the longitudinal field gradually increases to the E-field in the positive column.
- positive column:  
In this region the potential gradient is practically constant, and electron impact ionization occurs throughout this region.

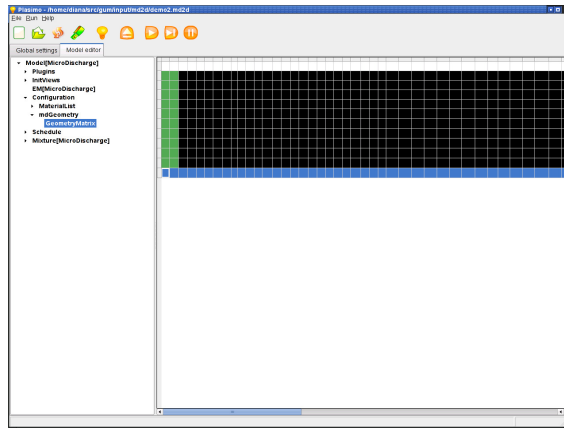
## Variation of the conditions

It is possible to change several parameters of the model to study the effects on the plasma. After every change it is necessary to install the model again, before it is run. The following changes can be applied:

- Geometry - a Cylindrical coordinate system.  
To change the Cartesian system into a Cylindrical coordinate system is relatively easy. First, in the **Model editor** tab you must expand the **Configuration** node in the tree by clicking the “+” symbol. Then select **mdGeometry**. On the right-hand side the geometry parameters will be shown, see Fig. 4.1(a). From the drop-down menu **Coordinate system** select **cylindrical** instead of **cartesian**.  
Note: When a cylindrical geometry is used axis of symmetry is located at the bottom of the geometry matrix. In Cartesian coordinates the bottom row in the matrix represents a dielectric. In cylindrical coordinates this row must be removed. To change this, open the subsection **GeometryMatrix** and simply select and delete (by right-clicking) the last row in the matrix. In order to be able to select a row or column, you must select “Grid layout mode” from the the right-click menu.
- Distance between the electrodes.  
To change the distance between the electrodes, or the radius of the discharge, you must only change **deltaX** or **deltaY** (Fig. 4.1(a)).  
As the anode is moved towards the cathode the positive column becomes shorter. However, when the anode is moved through the Faraday dark space the anode glow disappears, accompanied by a slight fall in the voltage necessary to sustain the discharge.
- Pressure.  
To change the pressure you must first select the **ModelEditor** tab, and then select **GasList**, which is located under the **Mixture** node.



(a) Geometry parameters



(b) Geometry Matrix

Figure 4.1: Accessing the geometry parameters and matrix.

How does the appearance of the discharge change with varying pressure?

An increase of pressure causes all the negative zones (cathode dark space, negative glow, and Faraday dark space) to be compressed towards the cathode. A decrease of pressure naturally causes the reverse effect: the negative zones expand and their boundaries become more diffuse. The positive column is driven into the anode and disappears altogether when the pressure is sufficiently low.

- Potential and current.

To define the potential as a function of time select `mdScheduleBlock` located in the `Schedule` node in the `Model Editor` tab. In the `Schedule` section time has been divided into ranges, so called “ScheduleBlock”’s, and for each of those one needs to specify the potential of each of the electrode materials. At the moment the order in which the different **Potentials** appear in the `ScheduleBlock` must match the order of the different electrode materials defined in the `MaterialList`!

If the potential is raised the length of the cathode dark space decreases, whereas the length of the negative glow increases slightly. In reality there is also an increase in the current through the discharge and consequently a general increase in the brightness of the luminous parts, since more energy is now being dissipated in the gas and some of this appears as additional radiation.

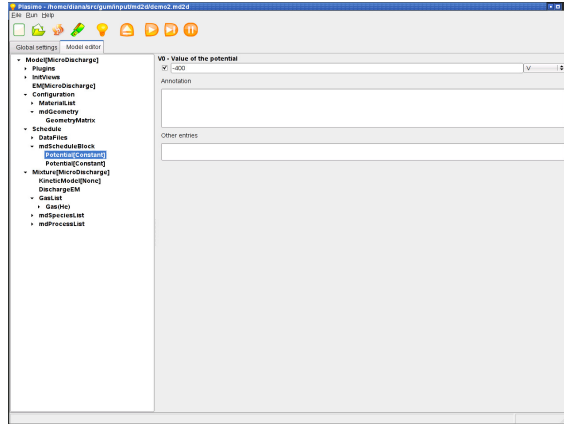


Figure 4.2: Setting the potential of the first electrode.

- Material of the cathode.

The cathode has a marked effect upon the voltage necessary to maintain the discharge. Lower voltages are needed when the cathode is a good emitter of electrons under bombardment by positive ions or photons. You can play with this by changing the secondary emission coefficients. The secondary emission coefficients are specified for each type of species, and are located in the **Model editor** tab under **Mixture / mdSpeciesList / mdParticle**

More input files are included in the **input** directory. One of these models is described in the following subsection.

### 4.3 MD2D sputtering model

**Example input file:** hcd-demo.md2d

**Location:** input/md2d/

The input file contains a model of a hollow cathode discharge (HCD) used for laser application. Description of the complete model you can find in [2, 3, 4], while here is a simplified version for demonstration purposes.

The geometry is a cylindrical hollow cathode with 4 mm inner diameter and 50 mm length; two anode rings are placed at both sides of the cathode, separated from each other with quartz rings. The geometry is symmetric around the center of the cathode (both axially and azimuthally).

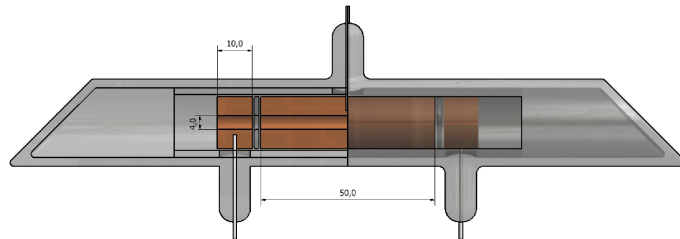


Figure 4.3: The hollow cathode geometry.

Hence, the simulated discharge region is half of the cathode length, the anode region and the dielectric in between. A fixed voltage of -300 V is applied.

The operating gas is He at constant pressure of 2.3 kPa. The considered species are  $\text{He}^+$ , Cu and  $\text{Cu}^+$ . The buffer gas atoms are ionized by electron impact ionization. The metal atoms in the discharge are produced only by sputtering due to ion bombardment of the cathode surface. The metal atoms are ionized in the plasma and they also can produce sputtering (self-sputtering). The ionization mechanism of Cu includes electron impact ionization and charge transfer with buffer gas ions.

## Sputtering module

The sputtering process can be taken into account in the model and it can be applied at different materials if the geometry consists many. This can be done:

- In section **Mixture** with the right mouse button add section **mdWallProcessList**
- In **mdWallProcessList** you must add the process **mdWallProcess** of type sputtering (the only available so far).
- Select **mdWallProcess**; on the right side window you can fill the species bombarding the wall (the projectile) and the sputtered atoms (recoils). You have to specify also the material index where the process should be applied. The indices are specified in the input file in section **Configuration/Material list**. In this example you see the cathode with index 1, the anode region with index 2, the dielectric with index 3 and index 0 is reserved always for the discharge region. The sputtering process is applied on materials with index 1 and 2.
- The sputtering yield for every projectile species is specified by selecting **mdSputteringYield**. The subsection **mdSputteringYield** appears automatically after adding **mdWallProcess** of type sputtering. The **mdSputteringYield** is of type **Constant**.

## 4.4 MC model

**Example input file:** `mc-demo.in`

**Location:** `input/mc/`

This model is simple kinetic drift model. We have Ar gas (density swarm) and we inject 10000 particles (electrons) into the simulation. The particles are injected as point sources at coordinates (0,0,0).

You can find the input file `mc-demo.in` in directory `input/mc/`.

After opening the input file, the main window will look like Fig. 4.4.

The main branches of the tree structure in the **Model Editor** tab are:

- **Environment:** this node contains the setup of the vessel, specifies the particles that are injected into the vessel, and defines the swarm list, i.e. the environmental particles the injected particles will collide with (the background gas).
- **Statistics:** obviously the necessary statistics are collected in this node, such as particle positions and velocities, and EDF's of the particle swarm that is followed. Moreover, the **Statistics** class can collect statistics of wall collisions (if there is any wall), such as position and velocity at the moment when the particle hits the wall, as well as EDF and deposition profile.
- **FlightControl:** this node controls the "flight" of the particles. Hence, it must know about all the reactions and the species involved in those reactions. For that reason the sections **mcSpeciesList**, **SwarmList**, and **mcProcessList** are set in the **FlightControl** section in the input file.

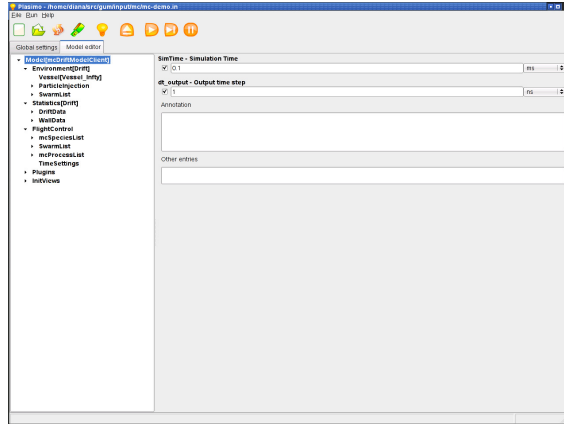


Figure 4.4: Main program window after loading the `mc-demo.in` input file.

After installing the model two more tabs appear:

- **General**: containing again the general common data;
- **MC Model Data**: containing all the output variables of the model.

### Variation of the conditions

As an exercise, the following changes can be applied to the model:

- Change the vessel. Put a plate at some distance from the injection point.
- Change the Electric field.
- Change the pressure (the density of the swarm).
- Inject more particles (or less).
- Change the Initial injection conditions.

## 4.5 PLASIMO model - 1

As an example of the plasimo module a model of an expanding DC plasma is treated in this section.

**Example input file:** `ArArc8_KG.gum`

**Location:** `input/demosuite/`

This model deals with a plasma created in an arc by means of a DC current. The plasma source has the shape of a cigarette; it is a cylinder with a radius of about 2 mm and a length of about 60 mm. The current creating this plasma is about 50 A. We write “about” because you can play with these plasma control parameters and look what effect it has on the plasma properties, like the electron density and temperature ( $n_e$ ,  $T_e$ ), the gas temperature ( $T_g$ ), and the flow field (velocities).

After loading the model, it can be explored by navigating through the leaves in the **Model editor**. Some of the leaves that are present are:

- **MainIterConfig:**  
This leaf contains some parameters to control the iterations.
- **PlasmaRegion:**  
This is the main part of the model, and contains a.o. the information of the grid and several fields.
- **TemperatureHP:**  
This leaf describes the heavy particle temperature. It contains a separate boundary condition (**BndConfig**) for every wall, and control of the iterations is handled in the **IterConfig** leaf.
- **TemperatureEL:**  
Similar to **TemperatureHP** only now for the electron temperature.
- **Mixture:**  
This describes the composition of the plasma, i.e. the particle species that are taken into account.

The characteristics of this plasma source are the same as that of a positive column (PC). For this PC a global model can be made, which will have the general outcome:

- The electron temperature is (almost) independent of the power (density), but is dictated by the electron loss frequency. The latter is often “diffusive”, meaning that this frequency more or less equals  $D_a/R^2$ , where  $D_a$  is the ambipolar diffusion coefficient, and R the plasma radius.
- The electron density ( $\text{m}^{-3}$ ) is determined by the power density ( $\text{Wm}^{-3}$ ).
- The heavy particle temperature is o.a. ruled by the electron density and the pressure (the plasma size...). So increasing  $n_e$  leads to higher  $T_h$ -values.

We will investigate these rules by running a Grand Model.

We are interested in the model for the expanding arc, named **ArArc8\_KG.gum**. This can be found in **input/demosuite**.

In the **EM[EMuniformECurrent]** node in the **Model editor** tab the current is set at 50 A. In our first run of the model we retain this (and all other) value(s); in exploring the plasma we can play with the current and select other values.

After installing the model three extra tabs appear:

- **General:** gives the status of the calculation and the convergence log graph. During run time the graph gives the residue as a function of the iteration number; right clicking on the graph makes it possible to select a.o. “X log scale” and “Y log scale”. Select the latter option.
- **Model regions**
- **Region data:** the left panel shows a tree structure containing a host of variables that are calculated by the model, organized in several groups: **Field(real-valued)**, **Field(vector)**, **Miscellaneous**, **Phi variable**, and **text**. The nodes under **Phi variable** are the transportables, thus the primary quantities of interest.

## Variation of the conditions

Make a table with columns headed by: **x1 velocity**, **HP temperature**, **Electron temperature**, **e-density**, and **Ohmic dissipation** (the latter quantity is not a transportable and can be found under **Field(real-valued)**). Now run the model for currents going from 50 A down to 5 A and fill in the table. Try to validate or falsificate the PC rules given above.



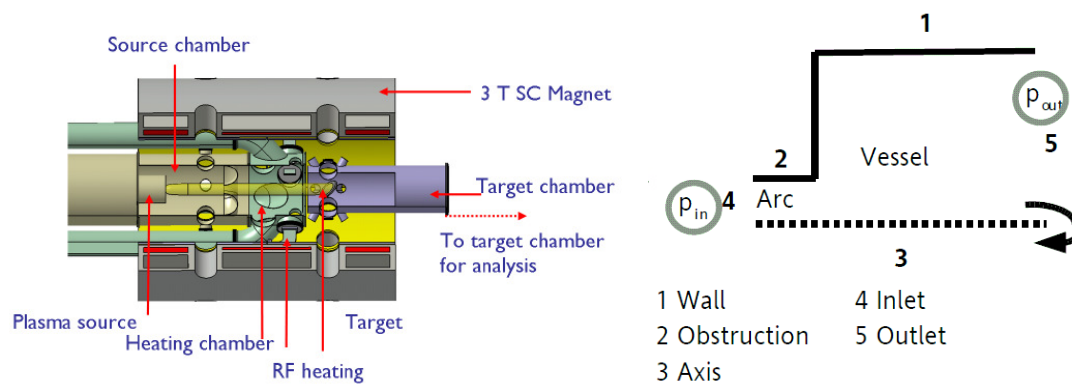
## 4.6 PLASIMO model - 2

A model of a transonic jet with multiple shock waves is treated in this section. Description of the complete model and detailed information you can find in [5].

**Example input file:** `transonicjet.gum`

**Location:** `input/demosuite/`

### Geometry



### Algorithm

- SIMPLE + Karki corrections
- Incompressible:  $\Delta p \rightarrow \Delta v$
- Compressible:  $\Delta p \rightarrow \Delta v / \Delta \rho$
- Karki corrections take into account influence  $p$  on  $\rho$

### Different flow regimes

The inlet and outlet pressure can be changed in the `Model editor` tab:

`Model/PlasmaRegion/Flow/Pressure/Function` (initialization)

`Model/PlasmaRegion/Flow/Pressure/BndConfig/BndCond` (boundary conditions)

The results can be seen in the `Model data` tab:

`Data/DefaultCylindricalGrid/Plasma/Barycentric`

`Data/DefaultCylindricalGrid/Plasma/Field(real-valued)/MachNumber`

- Set the Karki correction (`Model/PlasmaRegion/Flow/Pressure`) to 0.0: does the model still converge (`Data/General/Convergence log`)?
- What happens when you lower the inlet pressure; can you obtain a subsonic flow?
- Do subsonic flows converge without Karki corrections?

## 4.7 EM models

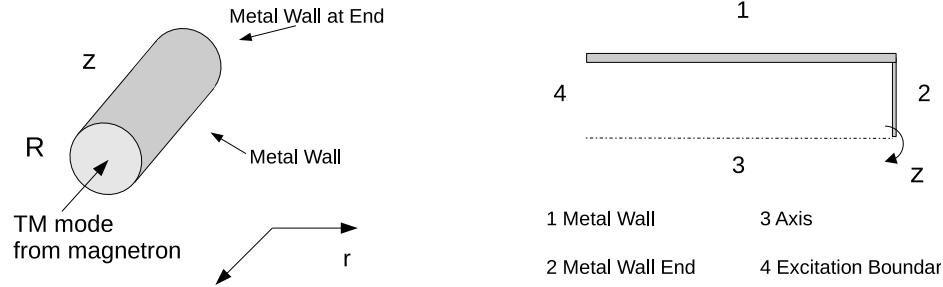
### 4.7.1 EM model - 1

**Example input file:** `circular_bessel.gum`

**Location:** `input/demosuite/`

The model represents transversal magnetic modes in a circular waveguide

#### Geometry



#### Algorithm

- Finite Differences in Frequency Domain
- Full Vector TM (Transversal Magnetic):  $E_r, E_z, \tilde{H}_\phi$

#### Standing waves and Evanescent Solutions

Install the program, and run it. Plot the normalized magnetic field  $\tilde{H}_\phi$

`Data/Cyl/Fields/em_region/Field (complex)/H_ud`

The excitation boundary creates a wave in  $z^+$ -direction, that interacts with the wave reflecting back from the metal wall at the end (2). Thus, standing waves are observed.

Below a critical radius,  $R_c$ , there are only evanescent (decaying) solutions instead standing waves. Decrease the radius  $R$ , until only evanescent solutions appear in the circular waveguide.

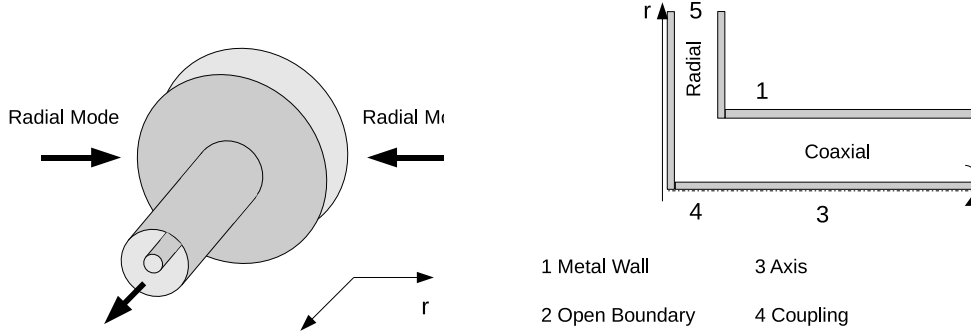
### 4.7.2 EM model - 2

**Example input file:** `radial_to_coaxial_wg_show.gum`

**Location:** `input/demosuite/`

The model represents radial-coaxial waveguide coupling.

## Geometry



## Algorithm

- Finite Differences in Frequency Domain
- Full Vector TM (Transversal Magnetic):  $E_r, E_z, \tilde{H}_\phi$

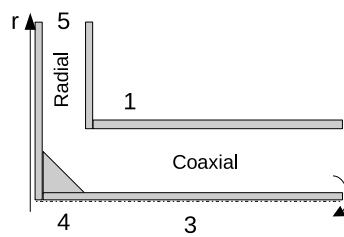
## Conversion from Radial to Coaxial solution

Install the program, and run it. Plot the normalized magnetic field  $\tilde{H}_\phi$

`Data/cyl/em_region/Field(complex)/H_ud`

The excitation boundary creates a wave in  $r^-$ -direction. When it reaches the coupling, the wave is converted into a wave propagating in  $z^+$ -direction. An open boundary condition (2) prevents most of the reflection, and almost no standing wave pattern is created in the coaxial waveguide. However, in the radial parallel plate waveguide, there is a standing pattern due to the reflection with the metal wire at the axis.

- Go to the Model editor and select in the hierarchial tree **Model/Grid/Geometry/CellMatrix**. You can now modify the geometry and draw a triangle (pyramidal cylinder in 3D), such as the one shown in the figure below.  
Right click and **Select Material**. Choose the **Metal**. Change the materials cells to metal by left clicking on them. Note that you can also enable the other section **CellMatrix**, where the triangle is already included.  
What happens to the standing waves in the radial waveguide?
- Change the boundary (2) from the open boundary condition (plane wave tangential E fields) to the homogeneous Dirichlet condition (null tangential E fields). You should notice a standing wave appearing in the coaxial waveguide.



- |                 |            |
|-----------------|------------|
| 1 Metal Wall    | 3 Axis     |
| 2 Open Boundary | 4 Coupling |

# Appendix A

## System requirements

the system requirements will be summarized for both Windows and Unix-like systems.

### A.0.1 Requirements for Microsoft Platforms

At present, native compilation of Plasimo on Microsoft Windows systems is not supported. Instead, Windows binaries are made available using cross-compilation on Linux, using the `mingw` cross-compiler suite, see <http://www.mingw.org/>. Please contact the Plasimo team at [info@plasimo.phys.tue.nl](mailto:info@plasimo.phys.tue.nl) if you want to obtain a binary version, suitable for Windows XP. No additional libraries are needed if you want to install Plasimo this way.

### A.0.2 Requirements for GNU/Linux and Unix Platforms

Table A.1 lists the packages which are needed to compile Plasimo on GNU/Linux and Unix systems. The end of this section gives detailed instructions on how to obtain these packages for the three most popular Linux distributions: Ubuntu, OpenSuse, and Fedora.

On most GNU/Linux installations you can check the availability of programs with the `which` utility. As an example, `which gcc` will print the location of the `gcc` program that is found in the path:

```
# which gcc
/usr/bin/gcc
```

If nothing is printed, the `gcc` program is not available and needs to be installed. Let us now assume that your system uses the `rpm` system to manage packages (like on Novell/SuSE, Redhat or Fedora systems). The version of the package that a file belongs to can be obtained as:

```
# rpm -qvf /usr/bin/gcc
gcc-4.1.0-25
```

In this case, `gcc` is present, the version being 4.1.0-25.

The remainder of this section explains the purpose of the various packages. You may wish to skip this information and go to section 2, the Quick Installation Guide instead.

In order to resolve some platform incompatibilities between the various flavours of Unix during the build process, use has been made of the so-called GNU build tools. These are the programs `gmake`, `aclocal`, `autoheader`, `automake`, `autoconf` and `libtool`. In addition, you need a C/C++ compiler. An overview of these packages is given in table A.1. The version requirements listed there are pessimistic: older versions of these products may work, but have not been tested with this version of the code. Note that `gmake` is also known as `make` on many GNU/Linux and Unix platforms.

`wxWidgets`<sup>1</sup> is a cross-platform toolkit for building graphical user-interfaces. On GNU/Linux and Unix only the GTK version has been tested, obviously this is our recommended version (see

---

<sup>1</sup>`wxWidgets` was formerly known as `wxWindows`.

Product	Version	Location
gcc	4.5.1	<a href="http://www.gnu.org/software/gcc/gcc.html">www.gnu.org/software/gcc/gcc.html</a>
automake	1.11.1	<a href="http://www.gnu.org/software/automake/automake.html">www.gnu.org/software/automake/automake.html</a>
autoconf	2.68	<a href="http://www.gnu.org/software/autoconf/autoconf.html">www.gnu.org/software/autoconf/autoconf.html</a>
libtool	2.2.6b	<a href="http://www.gnu.org/software/libtool/libtool.html">www.gnu.org/software/libtool/libtool.html</a>
gmake	3.82	<a href="http://www.gnu.org/software/make/make.html">www.gnu.org/software/make/make.html</a>
boost	1.42	<a href="http://www.boost.org">www.boost.org</a>
wxWidgets	2.8.11	<a href="http://www.wxwidgets.org">www.wxwidgets.org</a>
latex	3.141592-1.40.11-2.2	<a href="http://www.ctan.org">www.ctan.org</a>
latex2html	1.71	<a href="http://www.latex2html.org/">www.latex2html.org/</a>
pdflatex	3.141592-1.40-11-2.2	<a href="http://www.tug.org/applications/pdftex/">www.tug.org/applications/pdftex/</a>
doxygen	1.7.3	<a href="http://www.doxygen.org/">www.doxygen.org/</a>
dot	2.26.3	<a href="http://www.graphviz.org">www.graphviz.org</a>
python	2.7	<a href="http://www.python.org/">www.python.org/</a>
convert	6.6.5-8	<a href="http://www.imagemagick.org">www.imagemagick.org</a>

Table A.1: System requirements for building the software on GNU/Linux platforms. The products wxWidgets and below are weak requirements, the console versions of the Plasimo applications can be built without them. wxWidgets is required for graphical user-interface support (recommended), the remaining items are required for generating the documentation locally (see text).

the wxWidgets home page for details). Note that you need not only the libraries but also the development (header files) and that using wxWidgets may introduce additional software dependencies. We recommend to use the official packages of your favourite Linux distribution, if available.

The packages latex, latex2html, pdflatex, doxygen, dot, python and ImageMagick (more specifically, the program `convert` which is part of it) are weak requirements: these are needed to locally build the complete documentation. For released versions of the code the documentation can also be obtained from the Internet. See chapter “Generation of the Plasimo Documentation” of the programmer’s guide and the Plasimo website <http://plasimo.phys.tue.nl/generated-docs/> for details.

Some work has been spent to add support for other compilers than gcc, such as the Intel C++ compiler for Linux, the SGI MIPSpro compiler for IRIX and the Compaq/DEC CXX compiler for DEC Unix/Tru64 and Linux. In some case some special steps must be taken in the build process, this will be discussed in section B.2.2.

## Ubuntu

The following instructions have been tested for Ubuntu 13.10 a.k.a. Saucy. They also work in older versions (tested down to 10.04<sup>2</sup>). The required packages are listed in table A.0.2. The same instructions also work for Debian 7 a.k.a. Wheezy.

## OpenSuse

The required packages for version 13.1 are listed in table A.0.2. This has also been tested for previous releases down to 11.3.

Note, that it might be necessary to run the command `texhash` as root for the `html.sty` style file from the `latex2html` package to be found.

<sup>2</sup>Ubuntu 10.04 ships with boost 1.40. To install 1.42 add the repository `ppa:boost-latest/ppa` to your installation: `sudo add-apt-repository ppa:boost-latest/ppa`, update your package list and install `libboost1.42-dev`. If you are asked to remove version 1.40, answer yes.

cvs	libtool	automake
autoconf	build-essential	gcc
g++	gfortran	libboost-dev
texlive-latex-base	latex2html	doxygen
graphviz	python	imagemagick
libglu1-mesa-dev	liblapack-dev	gnuplot
transfig	libwxgtk2.8-dev	

Table A.2: Packages required for Ubuntu 10.04 through 13.10 and Debian 7 (approximately 1.5 GB).

cvs	gcc	gcc-c++
gcc-fortran	boost-devel	superlu
automake	autoconf	libtool
make	wxWidgets-devel	Mesa-devel
texlive-latex	texlive-appendix*	latex2html
gnuplot	doxygen	graphviz
python	ImageMagick	transfig
lapack-devel		

Table A.3: Packages required for OpenSuse 13.1 downto 11.3 (approximately 1 GB). The package denoted by \* is not required for release 12.2 and earlier.

## Fedora

The following instructions are identical for Fedora versions 13 through 20. The required packages are listed in table A.0.2.

cvs	gcc-c++	gcc-gfortran
boost-devel	lapack-devel	automake
autoconf	libtool	make
texlive-latex	texlive-a4wide*	texlive-epstopdf*
texlive-subfigure*	texlive-appendix*	latex2html
gnuplot	doxygen	graphviz
ImageMagick	python	gtk2-devel
mesa-libGLU-devel	transfig	wxGTK-devel

Table A.4: Packages required for Fedora versions 13 through 20 (approximately 1 GB). The packages denoted by \* are only required for release 18 and up.

## Appendix B

# Installation & Configuration

This section provide an in-depth discussion of the download and installation processes.

### B.1 Downloading Plasimo

#### B.1.1 CVS

In order to facilitate the code development process and the management of multiple versions of it the CVS<sup>1</sup> revision control system has been used. In order to obtain the sources you will need to install this program if you do not yet have it. This section will discuss how to configure cvs, and how to use it to obtain a version of choice of the sources.

The cvs program provides numerous additional features to retrieve old versions, inspect the development history of files, comparing different versions by date, release identifier et cetera. Check out the on-line CVS manual<sup>2</sup> for details.

#### Setting up CVS

Assuming you have read access rights, you can use cvs to obtain the sources from the *CVS repository* on the Plasimo server, **plasimo.phys.tue.nl**. Because the server has been configured to accept secure (encrypted) connections only, you also need *secure shell (ssh)*, which will normally be installed on Unix/Linux machines<sup>3</sup>.

The first step is to instruct your cvs client to use ssh. This can be done by setting the **CVS\_RSH** environment variable to **ssh**. How this should be done depends on the shell you are using. Bourne-shell (bash) users do **export CVS\_RSH=ssh**, while the C-shell (csh/tcsh) expects **setenv CVS\_RSH ssh**. Check the manual of your shell if you are in doubt. It is convenient to have these variables set automatically by your shell's initialisation script, otherwise you need to set them every time you open a shell to start using cvs.

#### Checking out the Files

Now that we are equipped to use cvs, we may do an initial checkout of the sources. Firstly, switch to the directory where you want to install the sources. Then issue the command

```
cvs -q -d :ext:name@plasimo:/usr/local/src/cvs checkout gum
```

It may be necessary that you provide the fully qualified name of the server instead of just **Plasimo**, namely **plasimo.phys.tue.nl**. Furthermore, replace **name** with your login name on **Plasimo**.

---

<sup>1</sup>see <http://www.cvshome.org/>

<sup>2</sup>see <http://cvsbook.red-bean.com/>

<sup>3</sup>When using Microsoft Windows a program such as TortoiseCVS ([www.tortoisecvs.org](http://www.tortoisecvs.org)) can be used, or even an environment such as Cygwin ([www.cygwin.com](http://www.cygwin.com)).



Please note that all commands and options under Unix are case-sensitive. This will create the directory `gum` in the current working directory. The option `-q` instructs `cvs` to report non-trivial output only. If your connection to the `cvs` server is via a modem, you should specify `-z3` before `-q` in order to save time by compressing all data traffic.

You can specify a set of default `cvs` options by creating a file `.cvsrc` in your home directory. A typical `.cvsrc` looks as follows:

```
cvs -z3 -q
update -d -P
diff -u
```

The meaning of the last two lines, which configure the defaults for the `update` and `diff` subcommands will become clear in section B.1.1.

By default, the subcommand `checkout` of `cvs` will give you the HEAD version of the sources. As explained in section 2 this is the branch on which active development takes place, the code here is not guaranteed to function properly, or even compile! Unless you have developer ambitions yourself, you should therefore check out one of the stable releases, probably the latest.

For getting such a stable release of the Plasimo sources, replace the part `checkout gum` in the `cvs` command above by `checkout -r REVISION gum`. Here `REVISION` should be replaced by the symbolic name of the version you want to fetch. As an example, the symbolic name `plasimo_4_0_0` corresponds to version 4.0.0 of Plasimo. These tag names can be found on the Manual and Source Code Documentation page on the website <http://plasimo.phys.tue.nl/>.

In case you have a version of the code already and want to install a second one, rename the `gum` directory to `gum-2.8.1`, say, before fetching the second version by entering another `cvs checkout` command.

## Updating and Switching to Another Version

Before updating to the latest version or switching to a different stable release, you may want to inspect the changes you have made locally. The reason is that these changes may conflict with those made by others and checked in. This can be achieved by changing to the `gum` source directory, and entering

```
cvs diff -u > mychanges.diff
```

This command writes the differences you have made locally to a file called `mychanges.diff`. Inspect this file to see if you made any important changes that you don't want to get lost. If this is the case, the best thing is to put the relevant files in a safe place, before continuing. In the file some lines may start with question marks. Unless you willingly added files locally you can ignore these lines. If you want to discard your changes, you now delete the locally modified files.

If you have the HEAD version of the Plasimo sources installed already, you can merge the latest version of the source and documentation files from the server with your local working copy by switching to the `gum` directory and typing

```
cvs update -d -P
```

The update-specific option `-d` indicates that newly created directories are checked out as well; `-P` means that files which are no longer in the repository are removed from your local `gum` directory<sup>4</sup>. In order to switch to a version other than HEAD you use the same command, but in addition you specify the version to check out using the `-r` option:

```
cvs update -d -P -r ReleaseID
```

Where `ReleaseID` is to be replaced with the symbolic name, like `PLASIMO_2_0`. Finally, if you are using a specific version of Plasimo, but want to switch to the HEAD version, use the option `-A` once with the update command:

---

<sup>4</sup>Because this is almost always what you want, you best add the line `update -d -P` to the file `.cvsrc` in your home directory. The same holds for the line `diff -u`.

```
cvs update -A
```

In following update commands the option `-A` can be omitted.

We end with an important note: after updating Plasimo, *always* check the `NEWS` file for changes that may affect you.

## B.2 Building the GUM binaries

### B.2.1 Compilation on Linux/Unix Platforms

Here it is assumed that your compiler and linker are supported by the GNU build toolchain (the programs `autoheader`, `autoconf`, `automake` and `libtool`). This is definitely the case if you have a GNU/Linux system and use the `gcc` compiler. If you have another platform or compiler some tweaking may be required. In that case please read the notes in the next section. Here's the complete build recipe:

1. Switch to the *top level source directory*, `gum`.
2. Execute `./makeconf.sh`. This shell script will execute various GNU build tools in the proper order. Some steps are repeated in subdirectories. This may take a while (in the order of a minute). You may get a warning telling you that you should add `m4` macros to `aclocal`, which you can safely ignore. If the script fails, you may need to call it from a Bourne shell by invoking `/bin/sh` or `bash` before calling the script.
3. Create a *build directory* and switch to it. This is where you are going to compile the libraries and application. You can have multiple build directories, and such directories can be added and removed at any time. It is best to give them names which characterize the configuration they will contain, like `linux-debug`, `MipsPro-optimized` or `win32-cross`.
4. From the *build directory*, run `configure`. This script is located in the top-level source directory `gum`. `Configure` accepts various options, some of these are general, others specific to the `gum` source package. You can run `configure` with `--help` as single argument to get a list of available switches, the `gum`-specific are listed below. The default settings are shown as is, if you want to override one or more of these defaults, you need to explicitly pass the options between brackets to `configure`.
  - `(--with-wx[=wx-bindir])`, `--without-wx`:  
If you enable this option, the Plasimo applications with a graphical user interface will be built. This requires a recent version of the `wxWidgets` toolkit, a cross-platform GUI-library. If no argument is provided, `configure` will look for a file `wx-config` in the directories in the `PATH` environment variable, and use that one. You may specify an alternative version of `wxWidgets` by providing the name of the directory in which the corresponding `wx-config` resides, as in `--with-wx=/usr/local/wx-HEAD/bin`.
  - `(--with-opengl)`, `--without-opengl`:  
Building with OpenGL support enables some Plasimo extensions, among which a variety of viewers for two- and three-dimensional data sets. Note that this option can only be enabled if you have OpenGL libraries and headers properly installed. It also implies usage of a version of `wxWidgets` with OpenGL extensions enabled. If you enable OpenGL, and the `configure` script finds the `Fame` library<sup>5</sup> on your system it is possible to create MPEG clips from OpenGL windows.
  - `--with-samples`, `(--without-samples)`:  
By default, all samples and tests will be built. If you disable this feature, you can still create the samples by switching to the sample directory of your choice and do this part of the build manually, as in `cd plspecies/samples/particle; gmake`

---

<sup>5</sup>see <http://fame.sourceforge.net/>

- (`--with-doc`), `--without-doc`:  
If you enable this option, the documentation is created. This is a time-consuming process, and requires some auxiliary programs (doxygen, graphviz). If you can afford it, read the Plasimo documentation online on our web site [plasimo.phys.tue.nl/plasimo-doc/](http://plasimo.phys.tue.nl/plasimo-doc/) (login required).
- (`--enable-static`), `--enable-shared`,  
`--disable-static`, (`--disable-shared`):  
The default setting is to build dynamic libraries only. Plasimo relies on shared library loading, so use this switch to change this behaviour only if you know what you are doing.
- `--enable-debug`, (`--disable-debug`):  
By default debugging information will be generated. This is useful during code development, but significantly increases the size of the resulting libraries and applications. If you don't know how to operate a debugger, you can safely disable this option.
- (`--enable-optimize`), `--disable-optimize`:  
By default, the code is compiled without optimizations. Production versions of the code should be compiled with optimizations enabled: it makes a huge difference in execution speed. Note that on most platforms optimization prevents debugging, so you might want to have two build trees, one with optimization for fast runs and one with debugging information but without optimization. GNU's gcc compiler suite allows to enable debugging optimized code, but note that inlining reduces the visibility of symbols from within the debugger.
- (`--enable-opt-arch`), `--disable-opt-arch`:  
Using this flag enables optimizations for the specific processor present on the machine where the code is built. This can improve the performance of the resulting code by a few percent, but may prevent the binaries to work on machines with a different (older) CPU.
- (`--enable-colors`), `--disable-colors`:  
When `Log(X)` messages are output, these may be colored according to their loglevel, red for 3, yellow for 2, green for 1 and bright white for 0. The application outputs escape sequences to set the colors, which work on most terminal emulations, basically everything that's compatible with a vt100 in this respect. Use the option to get more easily readable output if your terminal emulation supports it (both X11 xterm and Linux console do).
- (`--enable-smp`), `--disable-smp`:  
The TBCI numerics library used supports distributing large vector or matrix operations amongst several local CPUs by using POSIX threads. This speeds up operations for *large* objects on multiprocessor (SMP) machines. (For small data sizes, TBCI just does not make use of multithreading.) Use this flag to build with support for this; it can be controlled a run-time via the `Threads` or `MaxThread` configuration options.

Running `configure` takes some time, up to a couple of minutes. If `configure` fails, it will normally be due to an ill-formed list of options, and tell you how to correct. Typical cases are: missing required libraries, or attempting to enable OpenGL without `wxWidgets`. You may check `config.log` if you need more details.

5. The next step is to run `make`. You need GNU make, note that on some systems this is available under the name `gmake` instead. This will create the required libraries and the Plasimo main applications. Depending on the options which were passed to `configure`, the `wxWidgets`-based graphical front-end, the OpenGL extensions, the documentation and the executables in the samples subdirectories will be compiled as well. Note that a fresh build takes a considerable amount of time, perhaps an hour or more.

If you have updated to a different version of Plasimo (see B.1), you are safe if you repeat all the steps above. You may save some time, however:

- If you know that none of the `configure.in` files have changed, you can skip the `./makeconf.sh` step. In case of doubt, always run `./makeconf.sh` or `gmake -f Makefile.dist` in the gum source directory.
- If you want to recompile with the same configuration options as before, you don't need to create a new build directory, but you can use one of those that you created before. Change to your build directory (whether it's new or not).
- If you are sure that none of the `Makefile.am` files have changed, you can as well skip the `../configure` step. As above: In case of doubt rerun the `../configure` command.
- If you run `../configure` again in an already existing build directory, you probably want to use the same configuration as before. To this end `configure` has been made to emit the script `reconfigure` in the build directory. This will invoke `configure` with the same arguments as the last time it was called.
- Recompilation with `gmake` should require significantly less time than the original make, unless lots of files have changed or some of the headers that are included by almost any file have changed.

If you encounter weird errors after an update and a partial rebuild, switch to the build directory and enter:

```
make clean
make
```

The reason for the problems may have been an error in the dependency calculation. As a result some files may not have been rebuilt erroneously, these two commands guarantee that all files will be recompiled.

## B.2.2 Miscellaneous Compilers on Linux/Unix

Other compilers than `gcc` have been used to compile Plasimo on Unix-like systems, like DEC/Compaq CXX, Intel C++ and SGI MipsPro. The following remarks apply when using compilers other than `gcc` in order to build Plasimo.

When calling `configure`, you can specify the C-Compiler by setting the environment variable `CC` and the C++-Compiler by using the environment variable `CXX`. The linker is specified by the variable `LD`. If the compiler needs a certain flag to operate properly, you can include it in the variable as well.

During the configure process, test programs will be run which check for certain compiler features and bugs; unfortunately it's difficult to preview and/or work around all of these. Some bugs or features may be influenced by additional compiler flags. Some of them are already known to configure and automatically added to the compiler flags. Some may not and need to be added at compile time. Use the make options `EXTRA_CFLAGS`, `EXTRA_CXXFLAGS`, and `EXTRA_LDFLAGS` to add extra flags for the C compiler, the C++ compiler and the linker respectively. As a typical example, for some GNU/Linux systems you need `EXTRA_CXXFLAGS=-D_BSD_SOURCE` if you compile with `wxWidgets` enabled, because `wxWidgets` uses string functions that are only defined on BSD like systems (and the GNU system defines them only if `_BSD_SOURCE` is set. As a last resort, you may need to fix the `Makefiles` if your compiler is too different from the GNU compiler.

Here's a example for compilation on a Silicon machine running IRIX and using the MIPSpro compiler (version 7.3.1.1m). The text is based on an e-mail by Bart Hartgers, who figured this all out:

1. Go to your build directory.

2. Create the standard C++ headers that MIPSpro is missing by typing

```
mkdir cmissing
cd cmissing
python ../../util/generate_cpp_c_headers.py
cd ..
```

Obviously, you need the python<sup>6</sup> interpreter installed in order to have this work. Otherwise ask us for a tarball with the missing headers for MIPSpro.

3. Call configure with specifying Linker, C- and C++-Compiler.

```
LD="CC" CC="cc" CXX="CC -I'pwd'/cmissing" ../configure \
--enable-isoc99 --enable-optimize --disable-debug
```

Note: This syntax will only work on a Bourne shell.

4. Call a script to fix the Makefiles for MIPSpro.

```
../util/fix-Makefile-mipspro
```

5. Compile specifying some extra flags:

```
EXTRA_LDFLAGS='-elf -shared -lc' EXTRA_CXXFLAGS='-g -O2 -exceptions
-ptused -FE:template_in_elf_section' EXTRA_CFLAGS='-O2' gmake
```

You can omit the `-O2` in case you don't want your compiler to optimize the generated code. In case your compilation aborts with an error, you may still want to compile the rest by using `gmake -k` instead of `gmake`.

6. The MIPSpro compiler needs different flags for the executables in the end. So, go to the `app/` directory and do

```
gmake clean
EXTRA_LDFLAGS='-call_shared -lc' EXTRA_CXXFLAGS='-g -O2 -exceptions
-ptused -FE:template_in_elf_section' EXTRA_CFLAGS='-O2' gmake
```

7. The compiler seems to make a mess of functions to call with `plComplex` arguments. I can't really figure out what is going on exactly.

Bart Hartgers concluded his email with the following words: "Yes, this is awful. It was much more so to figure this out. Expect things to get better in the future as some of this knowledge may be built into the `autoconf/make` machinery."

---

<sup>6</sup>see <http://www.python.org/>

# Bibliography

- [1] Jan van Dijk, Kim Peerenboom, Manuel Jimenez, Diana Mihailova, and Joost van der Mullen. The plasma modelling toolkit plasimo. *Journal of Physics D: Applied Physics*, 42(19), 194012 (14pp), 2009.
- [2] D. Mihailova, M. Grozeva, G.J.M. Hagelaar, J. van Dijk, W.J.M. Brok, and J.J.A.M. van der Mullen. *J. Phys. D: Appl. Phys.*, 41, 245202, 2008.
- [3] D. Mihailova, J. van Dijk, M. Grozeva, G.J.M. Hagelaar, and J.J.A.M. van der Mullen. *J. Phys. D: Appl. Phys.*, 43, 145203, 2010.
- [4] D.B. Mihailova. PhD thesis, Eindhoven University of Technology, The Netherlands, 2010.
- [5] K S C Peerenboom, W J Goedheer, J van Dijk, and J J A M van der Mullen. Integral simulation of the creation and expansion of a transonic argon plasma. *Plasma Sources Science and Technology*, 19(2), 025009 (9pp), 2010.