PVMOS manual

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I. INTRODUCTION

This is (or rather will be as the current state of this document is far from finished) the manual for the Photo-Voltaic MOdule Simulator (PVMOS). PVMOS is an ordinary differential equation solver using finite-differences specifically designed to electrically model solar modules. For more information on how that works I refer the reader to []. The purpose of this document is to document how PVMOS is operated and installed. In the following sections I discuss in order, the installation, basic usage and operating principles and finally a detailed discussion of all available functions.

II. INSTALLATION INSTRUCTIONS

DISCLAMER:

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This program comes with ABSOLUTELY NO WARRANTY. This is free software, and you are welcome to redistribute under certain conditions. You should have received a copy of the GNU General Public License along with this program. If not, see http://www.gnu.org/licenses/.

PVMOS is is a command-line application written entirely in C. For the operation PVMOS depends on several libraries, most notably cholmod, for the solving of sparse linear systems. PVMOS has been tested on Linux and Windows systems. To install PVMOS you need to compile the source, for which a Makefile is provided. To install PVMOS you thus need to

- 1. Install PVMOS's dependencies
- 2. Edit the Makefile
- 3. Compile the code (type make)
- 4. test the executable

The performance of PVMOS is mostly determined by the performance of the sparse linear solver (i.e. cholmod). For an optimal performance an optimized BLAS library must be used (the reference BLAS is comparatively slow). For an optimized BLAS there are several options. One option is to use ATLAS which is available for all common CPU architectures and gives a very decent performance. On some architectures you can use Open-BLAS, which gives a very good performance (Open-BLAS is an actively developed fork of the now unmaintained GoTo BLAS). Some CPU manufacturers also publish their own optimized BLAS libraries for their CPU's (e.g. Intel and AMD). However, those libraries are usually provided under stricter conditions. For example Intel's Math Kernel Library is available commercially or under the condition that it is used for non-profit purposes (for that it provides a performance which was marginally better than Open BLAS in my tests).

III. BASIC USAGE AND OPERATING PRINCIPLES

The input for PVMOS is a plain text file with commands. To solve a problem PVMOS is typically called from the command line with as an argument the filename describing the problem. With these input files you can specify the geometry of your solar cell/module including the local properties such as electrode sheet resistances and solar cell

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properties. You can also specify which calculations you want PVMOS should perform and what data to save. A call to PVMOS from the command line looks like this:

pvmos [verbose-option] <input-file>

where [verbose-option] is an option which how much information PVMOS outputs to stdout, and <input-file> is the plain text input file. We first describe the mesh data structure in more detail in the next section.

More or less all functions implemented in PVMOS manipulate meshes in one way or another. To understand PVMOS one needs to understand PVMOS's mesh data-structure. A mesh in PVMOS consists of rectangular elements. For each element we store the coordinates of the lower left corner (x_1, y_1) and of the upper right corner (x_2, y_2) . Furthermore each node has a unique ID number. In order to save memory space we do not store the electrical properties for every element. Instead, we store the electrical parameters per "area". To this end a mesh contains a list of area data-structures which contain parameters such as the electrode sheet resistances and the solar cell properties. Each element now refers to a certain area by means of an area-ID number. Each area also has a name for easy referencing by the user. Finally, once you solve the potentials within a mesh for certain applied bias voltages, the mesh will also contain arrays with the potentials in the front and back contact for each element for each of the applied bias voltages.

In PVMOS we can define more than one mesh at the same time (this is useful as you can build meshes from several meshes, e.g. join two meshes for single cells to one mesh with two series connected cells). In order to reference one particular mesh, each mesh has a name. To reference an area within a mesh you can refer to <mesh-name>.<area-name>, i.e. the name of the mesh followed by a dot and the name of the area within that particular mesh. Sometimes we need to select elements in a mesh (for example when we want to assign a set of elements to a certain area). To this end each mesh has a list of selected elements. If you select elements in a mesh the list is occupied by the element ID's, after which you can do operations on the selected elements. Note that elements are selected on a per mesh basis.

The input file is parsed by PVMOS which sequentially processes the file. PVMOS provides functions to generate and manipulate meshes such that you can generate meshes describing the problem by a sequence of commands. To this end each mesh you define has a name to reference it by. The following section provides a command-reference.

IV. PVMOS COMMAND REFERENCE

Creating Meshes	
Keyword	Description
newmesh	Create a new, rectangular mesh. The command takes six arguments:
	1. x1, x-coordinate of the lower left corner
	2. y1, y-coordinate of the lower left corner
	3. x2, x-coordinate of the upper right corner
	4. y2, y-coordinate of the upper right corner
	5. Nx, Number of elements in x direction
	6. Ny, Number of elements in y direction
	7. mesh-name, Name of the new mesh
joinmesh	Create new mesh by joining two meshes. Make sure the meshes touch but do not overlap. The function takes offset values as input which allow you to "shift" the second mesh to align it to the first. The command takes 5 arguments:
	1. x_off , x-offset in coordinate system of the second mesh
	2. y_off, y-offset in coordinate system of the second mesh
	3. mesh1-name, Name of the first mesh
	4. mesh2-name, Name of the second mesh
	5. mesh3-name, Name of the resulting mesh

joinmesh_h

Create new mesh by joining two meshes. Make sure the meshes touch but do not overlap. The function takes a y-offset value as input which allow you to "shift" the second mesh in the y-direction to align it to the first. The x-offset value is the maximal x-value found in the first mesh. The command takes 4 arguments:

- 1. y_off, y-offset in coordinate system of the second mesh
- 2. mesh1-name, Name of the first mesh
- 3. mesh2-name, Name of the second mesh
- 4. mesh3-name, Name of the resulting mesh

joinmesh_v

Create new mesh by joining two meshes. Make sure the meshes touch but do not overlap. The function takes an x-offset value as input which allow you to "shift" the second mesh in the x-direction to align it to the first. The y-offset value is the maximal y-value found in the first mesh. The command takes 4 arguments:

- 1. x_off, x-offset in coordinate system of the second mesh
- 2. mesh1-name, Name of the first mesh
- 3. mesh2-name, Name of the second mesh
- 4. mesh3-name, Name of the resulting mesh

SELECTING ELEMENTS

Keyword select_rect

Description

Select a rectangular area in a mesh. The command takes five arguments:

- 1. x1, x-coordinate of the lower left corner of the selected rectangle
- 2. y1, y-coordinate of the lower left corner of the selected rectangle
- 3. x2, x-coordinate of the upper right corner of the selected rectangle
- 4. y2, y-coordinate of the upper right corner of the selected rectangle
- 5. mesh-name, Name of the mesh

select_circ

Select a circular area in a mesh. The command takes four arguments:

- 1. x_c, center x-coordinate of the selected circle
- 2. y_c, center y-coordinate of the selected circle
- 3. r, radius of the selected circle
- 4. mesh-name, Name of the mesh

select_poly

Select an area within a polygon-contour. In order to use this command you must first load a polygon from file with the load_poly command. The command takes one argument.

1. mesh-name, Name of the mesh

${\tt load_poly}$

Load a polygon from file. This command is used in conjunction with the select_poly command. The command takes one argument.

1. mesh-name, Name of the mesh

deselect

deselects a selection within a mesh. The command takes one argument.

1. mesh-name, Name of the mesh

Manually changing the mesh topology

Keyword split_x

Description

Split selected elements in x-direction. If no elements are selected, all elements are split. As the topology of the mesh changed all selected nodes in the mesh are un-selected after this command. The command takes one argument

1. mesh-name, Name of the mesh

split_y

Split selected elements in y-direction. If no elements are selected, all elements are split. As the topology of the mesh changed all selected nodes in the mesh are un-selected after this command. The command takes one argument

1. mesh-name, Name of the mesh

split_xy

Split selected elements in both x- and y-direction. If no elements are selected, all elements are split. As the topology of the mesh changed all selected nodes in the mesh are un-selected after this command. The command takes one argument

1. mesh-name, Name of the mesh

split_long

Split selected elements in thier longest direction. If no elements are selected, all elements are split. As the topology of the mesh changed all selected nodes in the mesh are un-selected after this command. The command takes one argument

1. mesh-name, Name of the mesh

split_coarse

Split selected elements until the node-edges are all smaller than a given length. If no elements are selected, all elements are split. As the topology of the mesh changed all selected nodes in the mesh are un-selected after this command. The command takes two arguments

- 1. mesh-name, Name of the mesh
- 2. 1, Maximum edge length

simplify

Attempt to simplify a mesh. If elements are selected they are un-selected as the topology of the mesh changed. The command takes one argument

1. mesh-name, Name of the mesh

SAVING AND LOADING MESHES

Keyword

Description

savemesh

Save a mesh to file in the PVMOS binary format, so it can be loaded again at a later time (see the loadmesh command). The command takes two arguments.

- 1. mesh-name. Name of themesh to be saved.
- 2. file-name, filename to save the mesh to.

loadmesh

Load a mesh saved to file in the PVMOS binary format (see the savemesh command). The command takes two arguments.

- 1. file-name, filename of the file containing the mesh data.
- 2. mesh-name, Name to assign to the loaded mesh

Element-wise export of data

Keyword

Description

printmesh

Export the mesh in a manner that is plottable with the gnuplot program (www.gnuplot.info/). The resulting plot draws the contour of each element in the mesh. The command takes two arguments:

- 1. mesh-name, Name of the mesh
- 2. file-name, filename to save the data in.

The output file will contain coordinates in columns. For each element the file contains the coordinates of the lower left- and the upper right corners empty line:

- x1 y1
- x2 y1
- x2 y2
- x1 y2

<empty line>

printconn

Print lateral connections in the electrodes in a format plottable with gnuplot (www.gnuplot.info/). When plotting the file (with vectors) a vector is drawn between the center of each element to the center of the adjacent elements to which it is connected. This routine may be useful when inspecting generated meshes. The command takes two arguments:

- 1. mesh-name, Name of the mesh
- 2. file-name, filename to save the data in.

The output is coordinates in columns. For each element the file contains the following data where xc, yc is the center of the current element and xca_i, yca_i is the center coordinate of the i-th adjacent

printarea

Print the geometry of the mesh which identifies each element and the area it belongs to. The fileformat is laid out such that it is plottable with a surface plot in gnuplot (www.gnuplot.info/). The command takes two arguments:

- 1. mesh-name, Name of the mesh
- 2. file-name, filename to save the data in.

The output file will contain data in columns. The file contains coordinates, the element ID and the corresponding area ID. Note that the parameters for each area can be exported with the printpars command. For each element it plots 2 times 2 data lines with an empty line inbetween. Between the data of two elements are two empty lines. This file is formatted such that when plotted with "splot" in gnuplot you can plot a surface for each element in the mesh, which allows you to see the areas in the defined geometry. For each element the following data is printed to the file:

printV

Print the front and back electrode element voltages for each stored solution. The output is formatted for gnuplot's splot command, such that a surface plot plots each element individually. The command takes two arguments:

- 1. mesh-name, Name of the mesh
- 2. file-name, filename to save the data in.

The output file will contain data in columns. The file contains coordinates followed by the potential in the positive and negative electrodes for each solution. For each element it plots 2 times 2 data lines with an empty line inbetween. Between the data of two elements are two empty lines. This file is formatted such that when plotted with "splot" in gnuplot you can plot a surface for each element in the mesh. For each element the following data is printed to the file:

printpar

Print a summary of the parameters per area, including both area-name and area-ID. The command takes two arguments:

- 1. mesh-name, Name of the mesh
- 2. file-name, filename to save the data in.

printmesh_sel

Same as printmesh except that it only exports a selected area specified by its lower left and upper right corners. The command takes six arguments:

- 1. mesh-name, Name of the mesh
- 2. x1, x-coordinate of the lower left corner of the selected rectangle
- 3. y1, y-coordinate of the lower left corner of the selected rectangle
- 4. x2, x-coordinate of the upper right corner of the selected rectangle
- 5. y2, y-coordinate of the upper right corner of the selected rectangle
- 6. file-name, filename to save the data in.

printconn_sel

Same as printconn except that it only exports a selected area specified by its lower left and upper right corners. The command takes six arguments:

- 1. mesh-name, Name of the mesh
- 2. x1, x-coordinate of the lower left corner of the selected rectangle
- 3. y1, y-coordinate of the lower left corner of the selected rectangle
- 4. x2, x-coordinate of the upper right corner of the selected rectangle
- 5. y2, y-coordinate of the upper right corner of the selected rectangle
- 6. file-name, filename to save the data in.

printarea_sel

Same as printarea except that it only exports a selected area specified by its lower left and upper right corners. The command takes six arguments:

- 1. mesh-name, Name of the mesh
- 2. x1, x-coordinate of the lower left corner of the selected rectangle
- 3. y1, y-coordinate of the lower left corner of the selected rectangle
- 4. x2, x-coordinate of the upper right corner of the selected rectangle
- 5. y2, y-coordinate of the upper right corner of the selected rectangle
- 6. file-name, filename to save the data in.

printV_sel

Same as printV except that it only exports a selected area specified by its lower left and upper right corners. The command takes six arguments:

- 1. mesh-name, Name of the mesh
- 2. x1, x-coordinate of the lower left corner of the selected rectangle
- 3. y1, y-coordinate of the lower left corner of the selected rectangle
- 4. x2, x-coordinate of the upper right corner of the selected rectangle
- 5. y2, y-coordinate of the upper right corner of the selected rectangle
- 6. file-name, filename to save the data in.

printIV

Export the IV characteristics of the device. Exports a file with two columns, the first contains all the simulated applied voltages and the second the corresponding total currents. The command takes two arguments:

- 1. mesh-name, Name of the mesh
- 2. file-name, filename to save the data in.

surfVplot

Export the front and back electrode voltages for a specific solution. Unlike the print-commands like printV the data is interpollated and mapped on a regular mesh. The command takes eight arguments:

- 1. mesh-name, Name of the mesh
- 2. x1, x-coordinate of the lower left corner of the selected rectangle
- 3. y1, y-coordinate of the lower left corner of the selected rectangle
- 4. x2, x-coordinate of the upper right corner of the selected rectangle
- 5. y2, y-coordinate of the upper right corner of the selected rectangle
- 6. Nx, Number of points in the regular mesh along the x-direction
- 7. Ny, Number of points in the regular mesh along the y-direction
- 8. Va, Applied voltage (if the sepcified voltage is not available the closest value will be taken
- 9. file-name, filename to save the data in.

surfPplot

Export the local power density for a specific solution. Just like in the surfVplot command the data is interpollated and mapped on a regular mesh. The command takes eight arguments:

- 1. mesh-name, Name of the mesh
- 2. x1, x-coordinate of the lower left corner of the selected rectangle
- 3. y1, y-coordinate of the lower left corner of the selected rectangle
- 4. x2, x-coordinate of the upper right corner of the selected rectangle
- 5. y2, y-coordinate of the upper right corner of the selected rectangle
- 6. Nx, Number of points in the regular mesh along the x-direction
- 7. Ny, Number of points in the regular mesh along the y-direction
- 8. Va, Applied voltage (if the sepcified voltage is not available the closest value will be taken
- 9. file-name, filename to save the data in.

Manipulating local properties

Keyword

Description

assign_properties

Assign nodes to a defined area. If no nodes are selected all nodes in the mesh are assigned to the specified area. The command takes one arguments:

1. area-name, Name of the area (<mesh-name>.<area-name>)

set_Rp

Set the electrode resistance for the positive electrode for a specified area. If the specified area does not exist it will be newly created. The command takes two arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. value, Sheet resistance value (Ω)

 set_Rn

Set the electrode resistance for the negative electrode for a specified area. If the specified area does not exist it will be newly created. The command takes two arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. value, Sheet resistance value (Ω)

set_Rpvn

Set the contact resistance between the positive electrode and the negative node for the applied voltage. The command takes two arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. value, Contact resistance (Ωcm^2)

set Rnvn

Set the contact resistance between the negative electrode and the negative node for the applied voltage. The command takes two arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. value, Contact resistance (Ω cm²)

set_Rpvp

Set the contact resistance between the positive electrode and the positive node for the applied voltage. The command takes two arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. value, Contact resistance (Ωcm^2)

set_Rnvp

Set the contact resistance between the negative electrode and the positive node for the applied voltage. The command takes two arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. value, Contact resistance (Ωcm^2)

set_JV

Specify a tabular data set to use as a JV characteristics. The command takes two arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. file-name, Name of a file containing two columns, voltage and current density (V, Acm⁻²)

set_2DJV

Specify a two-diode model for the JV characteristics. The command takes eight arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. J01, Saturation current density for the first diode (ideality factor one) (Acm⁻²)
- 3. J02, Saturation current density for the second diode (ideality factor two) (Acm^{-2})
- 4. Jph, Photo current density (Acm⁻²)
- 5. Rs, Series resistance (Ωcm^2)
- 6. Rsh, Shunt resistance (Ω cm²)
- 7. T, Temperature (K)
- 8. Eg, Band gap (eV)

set 1DJV

Specify a one-diode model for the JV characteristics. The command takes eight arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. J0, Saturation current density (Acm^{-2})
- 3. nid, Ideality factor
- 4. Jph, Photo current density (Acm^{-2})
- 5. Rs, Series resistance (Ωcm^2)
- 6. Rsh, Shunt resistance (Ωcm^2)
- 7. T, Temperature (K)
- 8. Eg, Band gap (eV)

set_R

Specify a resistance for the JV characteristics. The command takes two arguments:

- 1. area-name, Name of the area (<mesh-name>.<area-name>)
- 2. R, Resistance (Ωcm^2)

NUMERICAL SETTINGS

Keyword

set_SplitX

Description

It is sometimes useful to prevent the adaptive meshing algorithms from splitting certain nodes in x-or y-direction. This commands toggles the splitting of nodes in x-direction for a specified area (per default all nodes can be split in all directions). The command takes one argument:

1. area-name, Name of the area (<mesh-name>.<area-name>)

set_SplitY

It is sometimes useful to prevent the adaptive meshing algorithms from splitting certain nodes in xor y-direction. This commands toggles the splitting of nodes in y-direction for a specified area (per default all nodes can be split in all directions). The command takes one argument:

1. ${\tt area-name},$ Name of the area (<mesh-name>.<area-name>)

maxiter	Set the maximum number of iterations for solving the non-linear system. The command takes one argument:
	1. maxiter, Maximum number of iterations (default: 25)
${\tt tol}_{ extsf{-}}{\tt V}$	Absolute voltage tolerance for the break-off criterion. The command takes one argument:
	1. tol_V, Absolute voltage tolerance V (default: 10^{-5} V)
rel_tol_V	Relative voltage tolerance for the break-off criterion. The command takes one argument:
	1. tol_V, Relative voltage tolerance $-$ (default: 10^{-5})
tol_kcl	Absolute current tolerance for the break-off criterion (KCL stands for Kirchhoff's Current Law). The command takes one argument:
	1. tol_kcl, KCL tolerance A (default: 10^{-5} A)
rel_tol_kcl	Relative current tolerance for the break-off criterion (KCL stands for Kirchhoff's Current Law). The command takes one argument:
	1. tol_kcl, Relative KCL tolerance – (default: 10^{-5})
Solving	
Keyword	Description
solve	Solve the system (do a voltage sweep). The command takes four arguments:
	1. mesh-name, Name of the mesh
	2. V_start, Start voltage
	3. V_end, End voltage
	4. N_step, Number of voltage steps
$adaptive_solve$	Solve the system and adapt the mesh at one specified voltage. The command takes four arguments:
	1. mesh-name, Name of the mesh
	2. V_a, Applied voltage
	3. threshold, Ralative threshold for node splitting, a parameter between 0 and 1 that controls how agressive the mesh is adapted, where lower values lead to a more agressive mesh adaption (typical values between 0.3 and 0.5).
	4. N_step, Number adaptive meshing iterations
Verbosity levels	
Keyword	Description
out_quiet	Set verbosity to the minimum (only says something when it crashes). The command takes no arguments.
out_normal	Set verbosity to the normal level. The command takes no arguments.
${\tt out_verbose}$	Output additional data that may be interesting. The command takes no arguments.
out_degug	Output additional data that is only interesting for someone who is chasing bugs in the code. The command takes no arguments.

V. EXAMPLE:MONOLITHICALLY SERIES CONNECTED CELLS AND A DEFECT


```
\# x1 y1 x2 y2 Nx Ny name
newmesh 0.0 0.0 0.5 30 50 30 cell_a
newmesh 0.0 0.0 8e-3 30 1 30 p1
newmesh 0.0 0.0 12e-3 30 1 30 da
newmesh 0.0 0.0 5e-3 30 1 30 p2
newmesh 0.0 0.0 5e-3 30 1 30
                           рЗ
newmesh 0.0 0.0 8e-3 30 1 30 cp
newmesh 0.0 0.0 8e-3 30 1 30 cn
# A mesh consists of elements. The elements in a mesh belong to areas. When a mesh is
# initialized, all elements are part of the samedefault area. All areas that are part of a mesh
# must have a name. The name of the default area is the same as the name of the mesh when it is
# initialized. You can change the properties of an area by using "set" commands. You can also
# define new areas and assign elements to these new areas. More on that later. We first define
# the properties of the default areas for each mesh. To refer to a certain area you must specify
# the mesh and the area name like so: <mesh-name>.<area-name>
# Here we set the diode properties of the area cell_a in the mesh cell_a:
# mesh.area JO n Jph Rs Rsh Temp Eg
set_1DJV cell_a.cell_a 1e-8 1.5 0.03 1e-3 1e4 300 1.12
# here we set the sheet resistances for the same area, for the positive and negative electrodes
set_Rp cell_a.cell_a 20
set_Rn cell_a.cell_a 1
# here follow various areas for the various meshes
# set_R sets the resistance between front and back electrode, here we remove the contact
set_R p1.p1
            1e90
set_Rp p1.p1 1e9
set_Rn p1.p1 1
set_SplitX p1.p1 0
set_1DJV da.da 1e-8 1.5 0.03 1e-3 1e4 300 1.12
set_Rp da.da 20
set_Rn da.da 1
set_SplitX da.da 0
set_R p2.p2
             1e-5
set_Rp p2.p2 20
set_Rn p2.p2 1
set_SplitX p2.p2 0
set_R p3.p3
set_Rp p3.p3 1e-3
set_Rn p3.p3 1e9
set_SplitX p3.p3 0
# We have two contact strips. Within the strips we remove the solar cell (define it as a
# resistors with a very high resistance). Either the positive or the negative electrode is made
# very conductive, and connected to the positive applied voltage or OV, respectively.
# we do not split elements in this mesh in the x direction.
set_R cp.cp
             1e90
set_Rp cp.cp 1e-3
set_Rpvp cp.cp 1e-8
set_SplitX cp.cp 0
set_R cn.cn
set_Rn cn.cn 0.1
```

set_Rpvn cn.cn 1e-8
set_SplitX cn.cn 0

```
# Now we start to assemble the complete mesh out of the various parts. When joining meshes we
# will use the coordinate system of the first mesh. The coordinate system of the second mesh can
# be shifted by using an x and y offset value. Note that you have to take care not to make the
# meshes overlap I have not (yet) inplemented error checking for this! Here I will use
# jointmesh_h, which joins meshes horizontally and keeps track of the x-offset value for me
# (i.e. the x-offset valueis computed as the maximum x-value in the first mesh).
# xoff y_off mesh1 mesh2 mesh_out
joinmesh_h 0.0 cell_a p1 cell_p1
# 0.5+0.008=0.508
joinmesh_h 0.0 cell_p1 da cell_p1da
# 0.508+0.012=0.52
joinmesh_h 0.0 cell_p1da p2 cell_p1dap2
# 0.52+0.005=0.525
joinmesh_h 0.0 cell_p1dap2 da cell_p1dap2da
# 0.525+0.012=0.537
joinmesh_h 0.0 cell_p1dap2da p3 cell_p1dap2dap3
# 0.537+0.005=0.542
# Note that we now have many meshes defined:
# 2. p1
# 3. da
# 4. p3
# 5. p3
# 6. cp
# 7. cn
# 8. cell_p1
# 9. cell_p1da
# 10. cell_p1dap2
# 11. cell_p1dap2da
# 12. cell_p1dap2dap3
# The last mesh is cell stripe including dead area. We can easily series connect several cells
# by joining several instances of the last mesh in a row. Here we series connect five cells
joinmesh_h 0.0 cell_p1dap2dap3 cell_p1dap2dap3 2cells
# 0.542+0.542=1.084
joinmesh_h 0.0 2cells cell_p1dap2dap3 3cells
# 1.084+0.542=1.626
joinmesh_h 0.0 3cells cell_p1dap2dap3 4cells
# 1.626+0.542=2.168
joinmesh_h 0.0 4cells cell_p1dap2dap3 5cells
# 2.168+0.542=2.710
# Now we add the contacts. Our final mesh will be called: cells
joinmesh_h 0.0 cp 5cells cp5cells
# 8e-3+2.710=2.718
joinmesh_h 0.0 cp5cells cn cells
# 2.718+8e-3=2.726
# Now we have three perfectly homogeneous cells. That is boring so we add a shunt in the middle
# cell. First we create a new area. This should be part of the cells mesh. We can create a new
# area simply by starting to specify its properties, a new area will automatically be created.
# set JV cells.shunt shunt.dat
set_R cells.shunt 1e-3
```

set_Rp cells.shunt 20
set_Rn cells.shunt 1

```
# So far the creation of this area has no effect as no nodes are assigned to the area. However,
# before we assign nodes to this area we need a better resolution as the nodes in the cell area
# are now 0.01x1.0 cm^2, which is not sufficient to specify a small defect. I will add a defect
# at the coordinate (0.7,15). Refining the mesh will be done by selecting nodes and subsequently
# splitting the nodes. As the nodes are long in the y direction we first split the nodes only in
# the y-direction. The select_circ creates a circular selection, Only nodes whose center fall
# within the specified circle are selected. The circle is specified by its center coordinates
# (xc,yc) and the radius, r. The split commands automatically use the selected nodes. Beware
# that if no nodes are selected prior to giving a split command, all nodes in the mesh are split
xc yc r mesh
select_circ 1.5 15 4 cells
split_y cells
select_circ 1.5 15 3 cells
split_y cells
select_circ 1.5 15 2 cells
split_y cells
select_circ 1.5 15 1 cells
split_y cells
select_circ 1.5 15 0.1 cells
split_y cells
select_circ 1.5 15 0.07 cells
split_y cells
select_circ 1.5 15 0.06 cells
split_y cells
# Note that I gradually reduced the radius of the circle to get a denser and senser mesh around
# the coordinate (0.7,15). The nodes at this coordinate are now about 0.01x0.0078 cm^2 large. As
# the nodes are now close enough to quadratic we now split the nodes three times in both x- and
# y- direction to get a satifactory resolution with nodes of about 10m x 10m. Note that in a
# regular, equidistant mesh such node sizes would result in meshes of almost 50 milion nodes
# where the mesh here is about 20000 nodes large.
select_circ 1.5 15 0.05 cells
split_xy cells
```

Now we have at the location of the shunt nodes of about 0.001x0.001 cm^2, this suffices for # a defect with a radius of about 0.01 cm and up. As we already defined the shunt properties # of the area "shunt" we only need to assign nodes to this area. To this and we select the nodes # and then assign them (note if you do not select nodes before you do this, all the nodes in the # mesh will be assigned to the shunt area.

- $\mbox{\tt\#}$ Now we print some stuff so we can check whether we made no mistakes.
- # printmesh prints the mesh (i.e. contour of each node)
- $\mbox{\tt\#}$ printpars prints the name, id number and parameters for each area in a mesh

```
# printarea prints the meah and adds element id and area id for each element
# printconn prints the connections between nodes (verctors from center to center of a node
# (bi-directional)
printmesh cells mesh_initial.dat
printpars cells pars.dat
printarea cells areas.dat
printconn cells conn.dat
# Finaly do some simulations, we start by adapting the mesh a few iterations (5). The adapttive
# solve command expects as arguments:
# 1: the mesh to adapt
# 2: the applied voltage
# 3: adaption factor, af, (lower value means more nodes are adapted each iteration)
# 4: number of iterations for the mesh adaption
# We also output the refined mesh
# mesh V af iterations
adaptive_solve cells 2.4 0.3 5
printmesh cells mesh_refined.dat
# Now we simulate an IV curve, save it to IV.dat and plot the spatial potential distribution
\mbox{\tt\#} for the available simulation that is closest to 4.0 \mbox{\tt V}
solve cells -0.5 4.0 46
printIV cells IV.dat
```

surfVplot cells 0.008 14.0 2.718 16 300 400 4.0 thin_film_detail.dat surfVplot cells 0.008 0 2.718 30 300 400 4.0 thin_film_wholedevice.dat

printV cells checkpot.dat