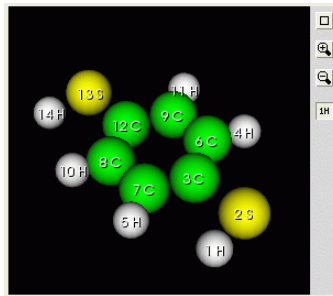


# Advanced Rappture Concepts

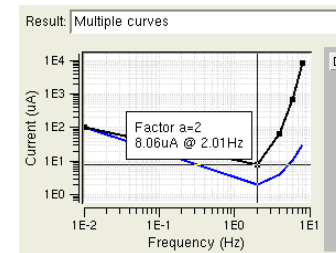


Carrier Statistics:

- Boltzmann
- Fermi**
- 2D Gas

Ambient temperature:

Grid points:



Michael McLennan  
Software Architect  
Network for Computational Nanotechnology

# Identify the elements

<group> of <group>'s

<choice>

<group>

<number>

<structure>

<box>

<field>

The screenshot shows the simulation interface with several elements highlighted in red boxes:

- Structure Materials Environment**: A tabbed menu at the top.
- Material: Si**: A dropdown menu showing the selected material.
- Minority carrier lifetimes**: A section containing two input fields: "For electrons: 1us" and "For holes: 1us".
- Structure**: A 3D diagram of a PN junction with three regions: P-type (purple), Intrinsic (white), and N-type (green).
- Doping (/cm<sup>3</sup>)**: A graph showing the doping concentration for each region. The P-type and N-type regions are both set to  $1e+17/cm^3$ , while the Intrinsic region is at 0.

Simulate new input parameters

## PN Junction Lab (v. 1.1padre)

Learn about any kind of P(I)N junction as you explore the devices in this simulator.

Input values for the various parameters on the left and click "Simulate" at the top to run the simulation. The parameters are currently set to model a standard PN junction diode. (no intrinsic region)

### - Material Properties

Define the material properties of the device, including elements and carrier lifetimes.

### - Structural Properties

Define the dimensional properties of the device, as well as the sample points taken along those dimensions.

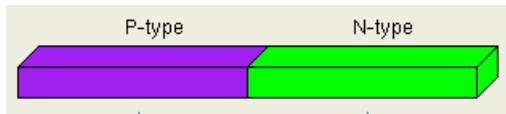
### - Temperature and Voltage

Set the ambient temperature and voltage sweep parameters.

### - Doping

Set the amount for doping for both P and N type materials. (Note: Intrinsic region always has zero doping)

## Structure of physical system being simulated

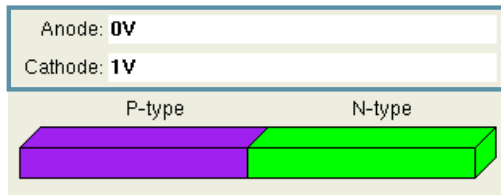


Just 1-D boxes,  
for now

```

<structure>
  <current>
    <components>
      <box>
        <about>
          <l a b e l >P-t y p e</l a b e l >< c o l o r >p u r p l e</c o l o r >
        </a b o u t >
        < c o r n e r >0</c o r n e r >
        < c o r n e r >0. 1 u m</c o r n e r >
      </b o x >
      <box>
        <about>
          <l a b e l >N-t y p e</l a b e l >< c o l o r >g r e e n</c o l o r >
        </a b o u t >
        < c o r n e r >0. 1 u m</c o r n e r >
        < c o r n e r >0. 2 u m</c o r n e r >
      </b o x >
    </c o m p o n e n t s >
  </c u r r e n t >
</s t r u c t u r e >
  
```

*Structure of physical system being simulated*

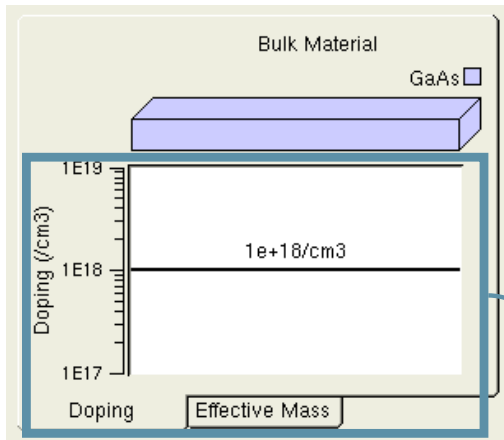


```

<structure>
  <current>
    <parameters>
      <number>
        <about><l a b e l >Anode: </l a b e l ></about>
        <u n i t s>V</u n i t s>
        <d e f a u l t>0V</d e f a u l t>
      </number>
      <number>
        <about><l a b e l >Cathode: </l a b e l ></about>
        <u n i t s>V</u n i t s>
        <d e f a u l t>1V</d e f a u l t>
      </number>
    </parameters>

    <components>...</components> ——— Same as before
  </current>
</structure>
  
```

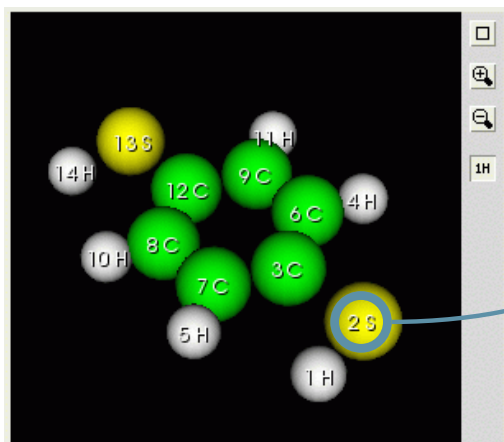
Structure of physical system being simulated



```

<structure>
  <current>
    <uni ts>um</uni ts>
    <parameters>
      <number i d="dopi ng" >...</number>
    </parameters>
    <components>...</components>
    <fi el ds>
      <fi el d>
        <about>
          <l abel >Dopi ng</l abel >
          <col or>bl ack</col or>
          <scal e>l og</scal e>
        </about>
        <uni ts>/cm3</uni ts>
        <component>
          <constant>dopi ng</constant>
          <domai n>box0</domai n>
        </component>
      </fi el d>
    ...
  
```

*Structure of physical system being simulated*

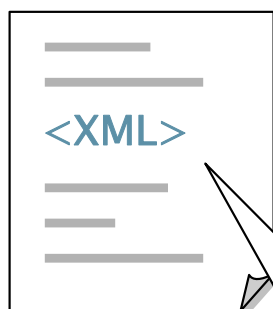


Turns atom labels on by default

```
<structure>
  <current>
    <components>
      <mol ecul e>
        <about><embl ems>on</embl ems></about>
        <formul a>pdt</formul a>
        <atom i d="0">
          <symbol >H</symbol >
          <xyz>-1. 24935  -3. 41562  0. 0</xyz>
        </atom>
        <atom i d="1">
          <symbol >S</symbol >
          <xyz>0. 08092  -3. 19426  0. 0</xyz>
        </atom>
        ...
      </mol ecul e>
    </components>
  </current>
</structure>
```

# Focus on <output> side of tool.xml

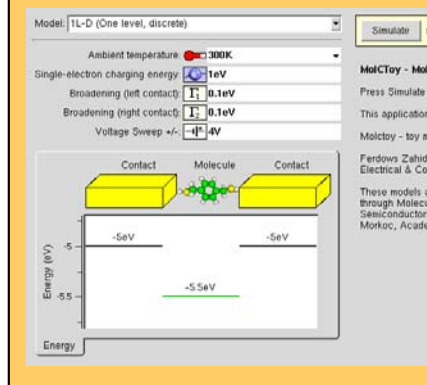
description of tool,  
including inputs  
and outputs



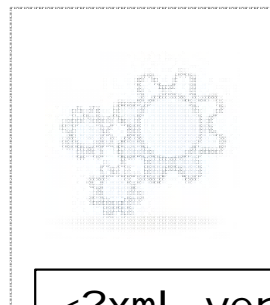
tool.xml



Rappture



Produces the  
user interface  
*automatically!*



```
<?xml version="1.0"?>
<run>
  <tool >
    <about>This is my tool.</about>
    ...
  </tool >
  <i nput >
    [Red dashed box]
  </i nput >
  <output >
    [Green solid box]
  </output >
</run>
```

## Standard output from simulator

```
Result: Output Log
***** ADEPT/F - 2.1   input file: adp20638

 1 *title   input generated by adeptwr

 2 mesh    nx=250 xres=0.5
 3 misc    tempk=300

 4 * layer Al(0.3)Ga(0.7)As
 5 layer   tm=0.2 nd=1e+17
 6 +      eg=1.797 chi=3.827 ks=11.9
 7 +      nc=8.57e+17 nv=1.11e+19
```

Controls for search through text

Treated as unimportant  
(low level) output, and  
therefore listed last

```
<output>
  <l og>***** ADEPT/F - 2.1   input
file: adp20638       Sat Jul 30 19:39:36
2005 *****

 1 *title   input generated by adeptwr

 2 mesh    nx=250 xres=0.5
 3 mi sc   tempk=300

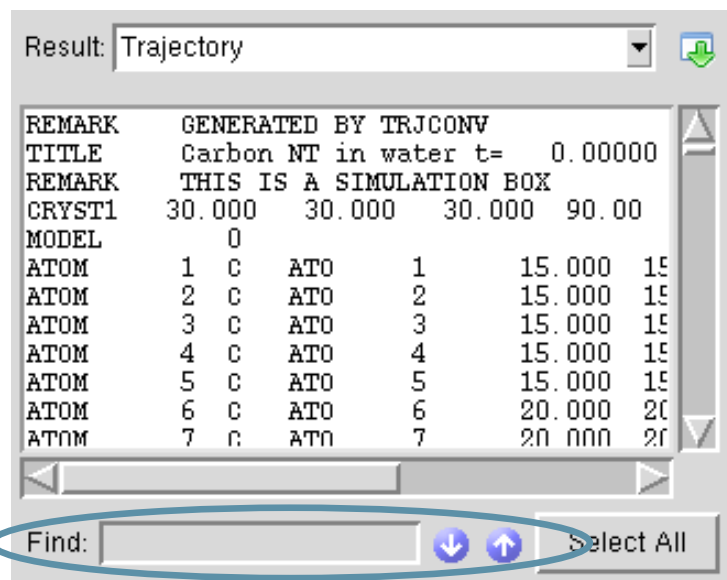
...
</l og>
</output>
```

or, in Python...

```
import Rappture
import sys
driver = Rappture.Library(sys.argv[1])
...
driver.put('output.log', stdout)
```



Other output files from simulator—including binary files

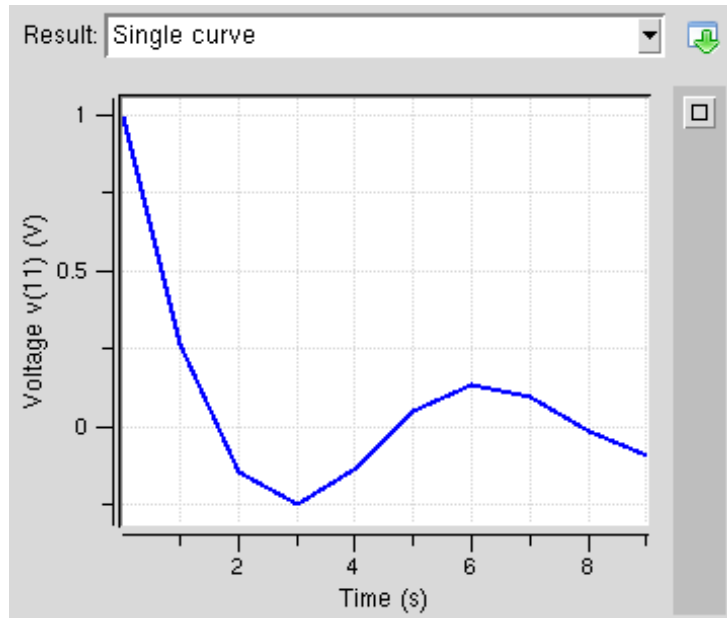


Controls for search through text

```
import Rappture
import sys
driver = Rappture.Library(sys.argv[1])
...
path = 'output.string(traj)'
```

```
<output>
  <string id="traj">
    <about>
      <label>Trajectory Data</label>
      <description>Data in pdb
    </description>
    </about>
    <current>REMARK      GENERATED BY
TRJCONV
TITLE      Carbon NT in water t=  0.00000
...
    </current>
  </string>
</output>
```

## X-Y plots



```

import Rappture
import sys
driver = Rappture.Library(sys.argv[1])
...
path = 'output.curve(single)'

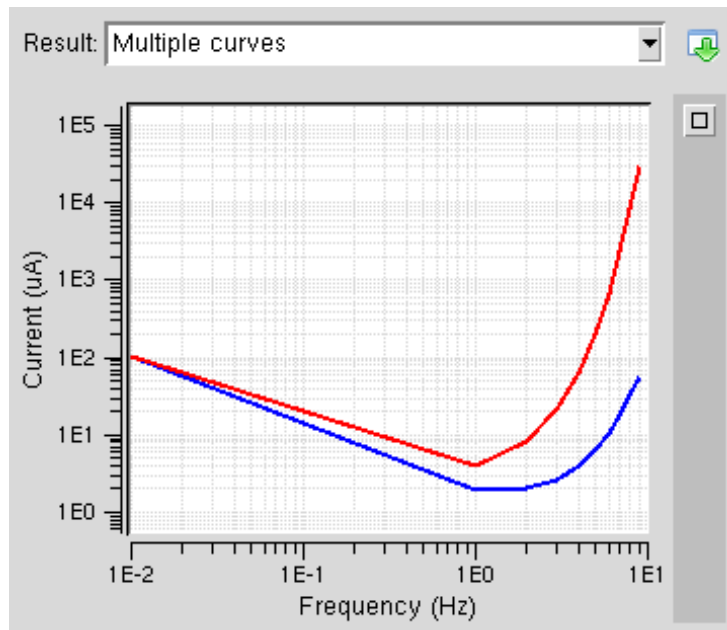
driver.put(path+'.about.label',
           'Single curve')

driver.put(path+'.xaxis.label', 'Time')
driver.put(path+'.xaxis.units', 's')
driver.put(path+'.yaxis.label', 'Voltage')
driver.put(path+'.yaxis.units', 'V')

data = ""
0 0
1 2
3 4 ""
driver.put(path+'.component.xy', data)

```

## Multiple curves on the same plot

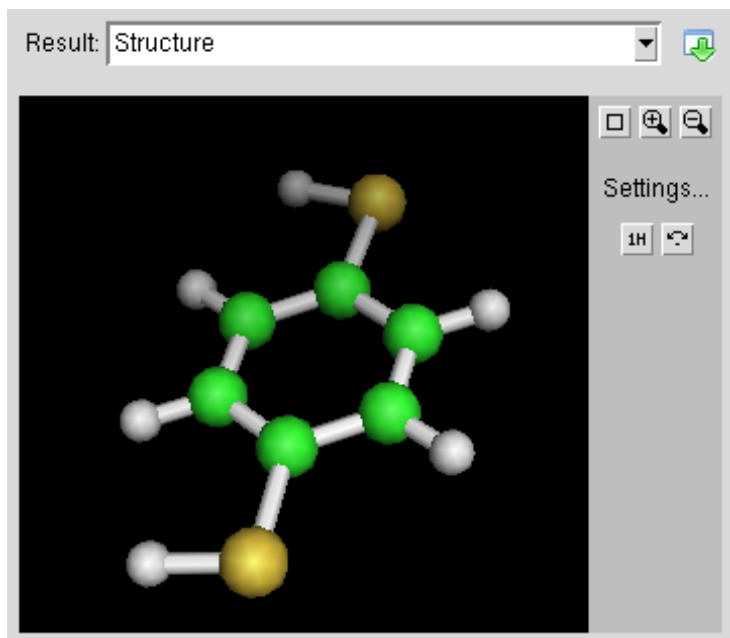


Different labels  
for different curves

```
import Rappture
import sys
driver = Rappture.Library(sys.argv[1])
...
path = 'output.curve(m1)'
driver.put(path+'.about.group',
           Multiple curves')
driver.put(path+'.about.label',
           factor a=1)
...
path = 'output.curve(m2)'
driver.put(path+'.about.group',
           Multiple curves')
driver.put(path+'.about.label',
           factor a=2)
```

Same group name

## Molecules



```

import Rappture
import sys
driver = Rappture.Library(sys.argv[1])
...
path = 'output.structure(mol)'
driver.put(path+'.about.label', 'Structure')

path += 'components.molecule'
driver.put(path+'.atom(0).symbol', 'H')
driver.put(path+'.atom(0).xyz', xyz0)
driver.put(path+'.atom(1).symbol', 'S')
driver.put(path+'.atom(1).xyz', xyz1)
...

```

*Scalar field defined over (x,y) or (x,y,z)*

```

Result: 3D Wavefunctions
import Rappture
object 1 class gridpositions counts 126 30 22
origin 0.00000000E+00 0.00000000E+00 nx ny nz 00E+00
delta 50.0 0.0 0.0 xgrid
delta 0.0 50.0 0.0 ygrid
delta 0.0 0.0 69.0476190476 zgrid
object 2 class gridconnections counts 126 30 22
object 3 class array type double rank 0 items 83160 data follows
0.28865594E-02
0.28865594E-02
0.28865594E-02
...
-0.71134413E-02
-0.71134413E-02
attribute "dep" string "positions"
object "regular positions regular connections" class field
component "positions" value 1
component "connections" value 2
component "data" value 3
    
```

Scalar values:  
z-index varies fastest,  
then y-index,  
then x-index

total number of  
data points:  
nx \* ny \* nz

...[1])  
(k', dxdata)  
type',  
city 1')  
(k', dxdat2)  
type',  
-max 1')

## Sequence of images, curves, or fields



```

<sequence id="movie">
  <about>
    <label>Animated sequence</label>
  </about>
  <index><label>Frame</label></index>

  <element id="0">
    <index>1</index>
    <image>
      <current>/9j/4AAQSkZJRgAA...</current>
    </image>
  </element>

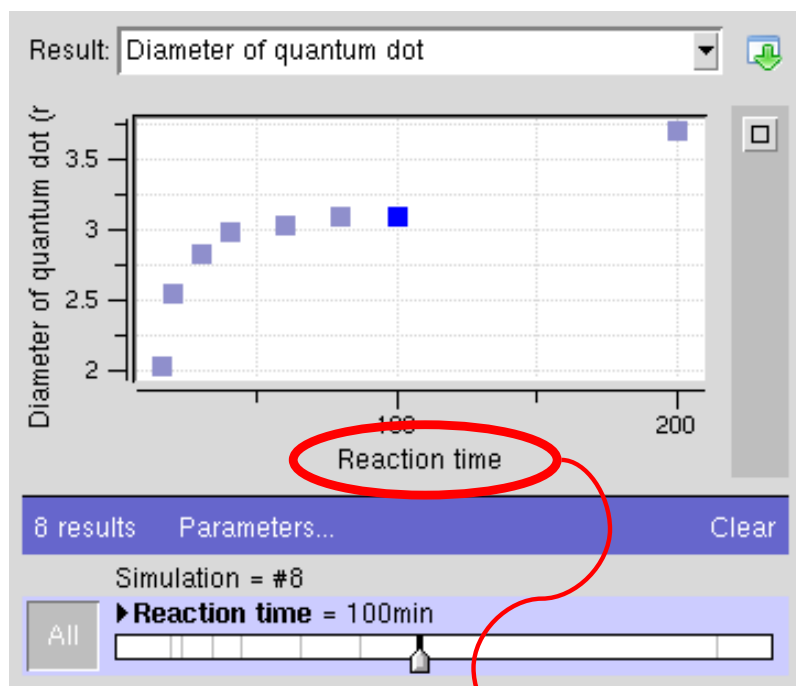
  <element id="1">
    <index>2</index>
    <image>
      <current>/9j/4ARgASkZJQQR...</current>
    </image>
  </element>

  ...
</sequence>
  
```

Just like a normal  
output image

## <number> and <integer>

*Just a number or integer, but compare across many runs*



Axis changed

```
import Rappture
import sys
driver = Rappture.Library(sys.argv[1])
...
path = 'output.number(d)'

driver.put(path+'.about.label',
           'Diameter of quantum dot')

driver.put(path+'.units', 'nm')
driver.put(path+'.current', d)
...
```

rappture\_xml\_elements - Rappture - Trac - Mozilla Firefox

https://developer.nanohub.org/projects/rappture/wiki/rappture\_xml\_elements

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## Zoo of Inputs/Outputs

Welcome to the zoo of Rappture elements! This page shows the various snippets of XML code needed to create a Rappture tool.xml file.

### Overall Structure

Each tool is described by a tool.xml file, which has the following structure:

```
<?xml version="1.0"?>
<root>
  <tool>
    <title>Name of the tool</title>
    <about>Description and credits</about>
    <command>
      @tool/path/to/executable @driver
    </command>
    <limits>
      <runtime>900</runtime>
      <filesize>1000000</filesize>
    </limits>
    <layout>xxx</layout>
    <control>xxx</control>
    <analyser>xxx</analyser>
    <reportJobFailures>1</reportJobFailures>
  </tool>
  <input>
    ..see Element Index below...
  </input>
  <output>
    ..see Element Index below...
  </output>
</root>
```

The <tool> section describes the underlying compute engine and includes the command needed to run it. This can be any Unix-style command line. The @tool keyword gets replaced with the name of the directory containing the tool.xml file. The

Carrier Statistics: Fermi  
Boltzmann  
Fermi  
2D Gas

Ambient temperature: 300K

Grid points: 100

## Zoo of Examples

- Complete catalog of data objects online
- See screen shots
- Copy xml code