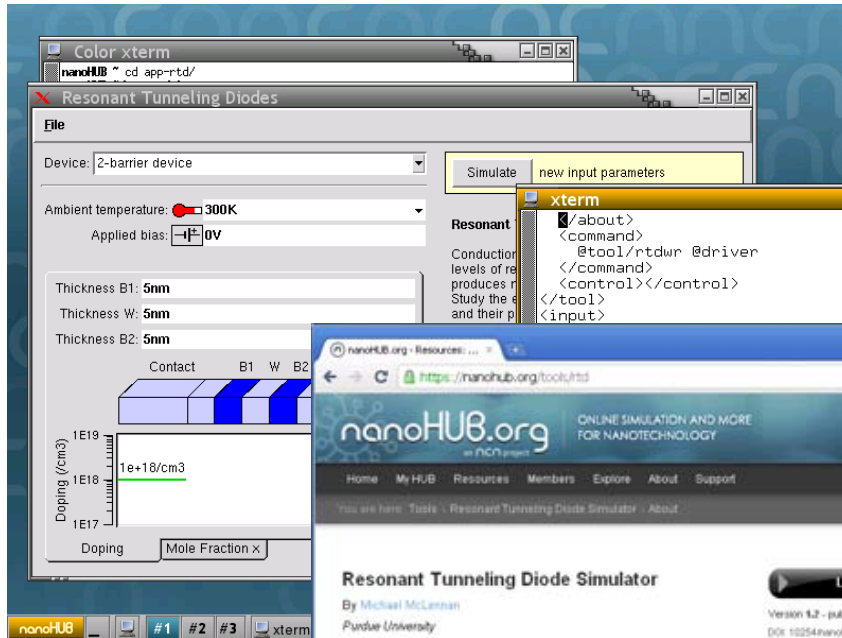


Regression Testing

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Build the tool...



Install, approve...

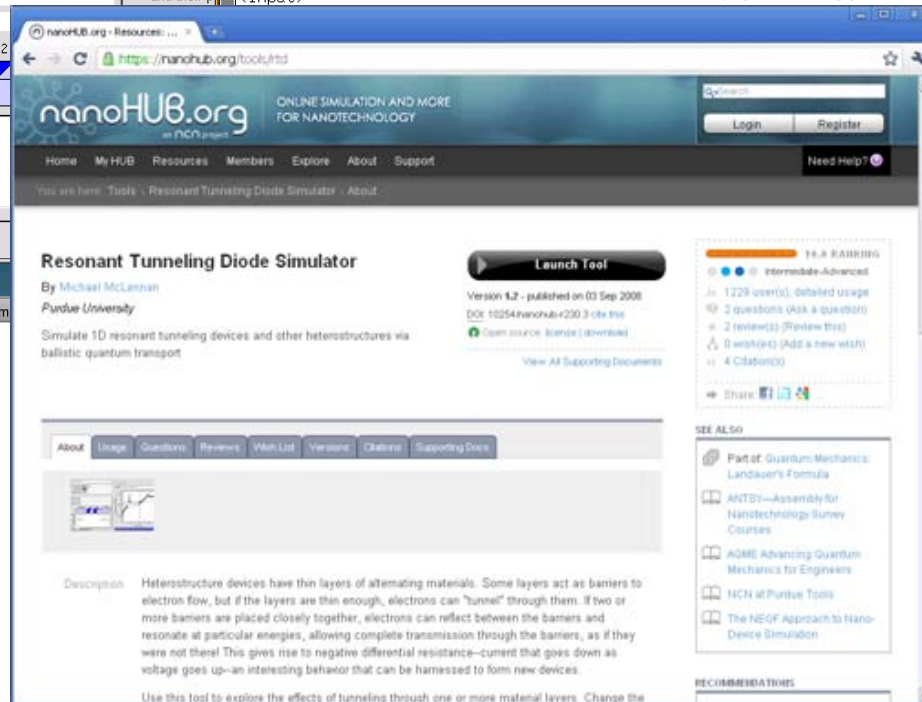
What's next?

Your latest code is installed and ready on nanoHUB.org. Please test your tool by clicking the button below to make sure that everything is working properly, as well as verify that the page describing your tool is created and displays correct information:

- Test your application:

Launch tool →

- Review the page describing your tool



is working properly, click here to

ed in your latest fixes, click here to

the latest version for testing and

Published!

Usually doesn't end there...

-
- Allow download of 3d simulation data

+7 0 +5
 - 257439 When I launch this tool, it crashes right away and collapses down to a...

open
-
- Proposed by Golam Rabbani on 31 May 2009, 2 Comments
-
- by Michael McLennan ([mmc](#))
-

Wishes

Bugs

New physics

$$\frac{m^* v_{emit}^2}{2} \approx \hbar \omega_{phonon(opt.)} \quad \mu = m_{sat}^2 \frac{2L}{W} \frac{1}{C_i}$$

```

Color xterm
real hbar, hbar2, Kb, m0, eVtoJ, pi, q
real ks, mtocm
parameter ( hbar=6.58216e-16,
+          hbar2=hbar**2,
+          Kb=8.61738e-5,
+          m0=9.1095e-31,
+          eVtoJ=1.60218e-19,
+          pi=3.141592654,
+          q=eVtoJ,
+          ks=12.846,
+          mtocm=100.0 )
c
c DEFINITIONS --
c ltor ... electrons traveling left to right
c rtol ... electrons traveling right to left
    
```

What's next?

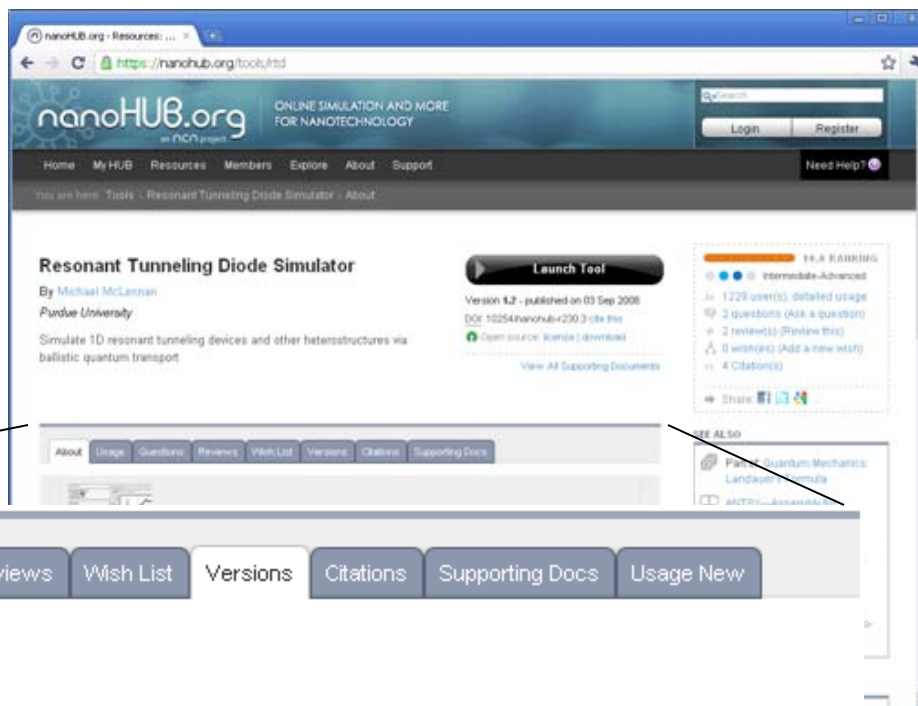
Your tool has been published and is now open to the public. Users will find out about your tool by reading the following page. Be sure to keep this information up to date:

<http://nanoHUB.org/tools/rtd>

Your options:

- › I've made changes [Please install the latest code for testing and approval.](#)





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Versions

Version	Released	DOI Handle	Published
3.01	03 Dec 2010	10254/nanohub-r1307.7	✓
3.0	03 Nov 2010	10254/nanohub-r1307.6	✗
2.1	17 Jul 2009	10254/nanohub-r1307.5	✓
2.02	25 Sep 2008	10254/nanohub-r1307.4	✗
2.0.1	08 May 2008	10254/nanohub-r1307.3	✗

You can publish multiple versions and keep more than one active

- Build up a suite of test cases
- Run them as you go along to make sure they still work
- Investigate failures and track down the problem

The screenshot shows the Rappture Regression Tester window. The left pane displays a tree view of test cases with their results:

Result	Test Case
✗	helium
✗	2eV
✗	1eV
✗	0eV
✗	fail
	multiple
	levels
✓	spaces are allowed...
	roomtemp
✗	1V
✗	0eV
	nitrogen
✓	1eV
✓	0eV

The right pane shows details for the selected test case '1V':

Test: 1V
Test failed

This test should generate a warning due to differences in the input parameters.

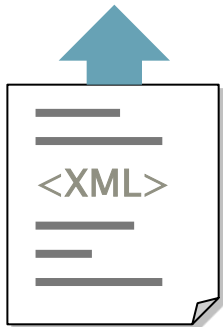
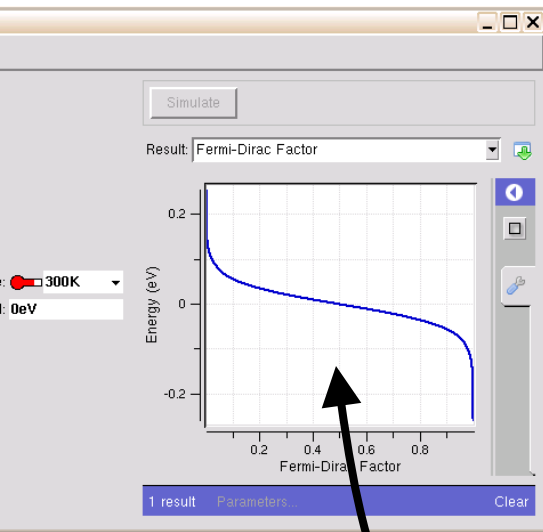
Differences:

- ⚠ **Input: Control value**
input.boolean(extra)
Test case has this extra input value
- ✗ **Output: Extra number in output**
output.number(extra)
Result is missing from current output
- ✗ **Output: Fermi-Dirac Factor**
output.curve(f12)
Result differs from expected value

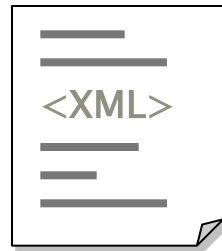
Buttons: Run, << New golden standard

1 Run the desired test case...

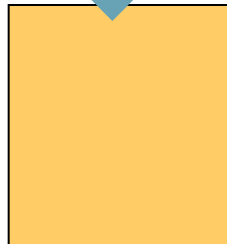
Runner



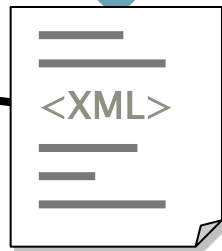
tool.xml



driver1



Your Program



run12703129102.xml

2 Move into tests directory

```

Color xterm
$ ls
docs  spiro.m  tool.xml
$ mkdir tests
$ unset SESSIONDIR
$ rapture
$ ls
docs  run12703129102.xml  spiro.m  tes
$ mv run12703129102.xml tests
$ █
    
```

Results are moved out of the current working directory to the "results" directory **unless you unset SESSIONDIR**



~/data/results/\$SESSION

3 Add label/description to each test

```
Color xterm  
$ vi tests/run12703129102.xml
```

```
<?xml version="1.0"?>  
<run>  
  <test>  
    <label>roomtemp|0eV</label>  
    <description>Should work at room temperature and 0eV.</description>  
  </test>  
  <tool>  
    <about>Press Simulate to view results.</about>  
    <command>tclsh @tool/fermi.tcl @driver</command>  
  </tool>  
  <input>  
    <number id="temperature">  
      <about>  
        <label>Ambient temperature</label>  
        <description>Temperature of the environment.</description>  
      </about>
```

separator for test folders

X Rappture Regression Tester

Result	Test Case
	helium
✖	2eV
✖	1eV
✖	0eV
✖	fail
	multiple
	levels
✓	spaces are allowed...
	roomtemp
✖	1V
✖	0eV
	nitrogen
✓	1eV
✓	0eV

Select: All None

Test: 1V
Test failed

This test should generate a warning due to differences in the input parameters.

Differences:

- ⚠ **Input: Control value**
input.boolean(extra)
Test case has this extra input value
- ✖ **Output: Extra number in output**
output.number(extra)
Result is missing from current output
- ✖ **Output: Fermi-Dirac Factor**
output.curve(f12)
Result differs from expected value

Run

<< New golden standard

Output results:

- ✘ Output value has changed
- ✘ Output value is missing
- ✘ Output value is extra (not supposed to be there)

Input values:

- ✘ Input value has changed--label, units, etc.
- ✘ Input value is missing from test case
- ✘ Input value is extra (not supposed to be there)

Fix the tool

```

Color xterm
real hbar, hbar2, kb, n0, eVtoJ, pi, q
real ks, ntcoc
parameter ( hbar=6.58216e-16,
+          hbar2=hbar**2,
+          kb=9.61739e-5,
+          n0=9.1095e-31,
+          eVtoJ=1.60218e-19,
+          pi=3.141592654,
+          q=eVtoJ,
+          ks=12.846,
+          ntcoc=100.0 )
c
c DEFINITIONS --
c ltor ... electrons traveling left to right
c rtor ... electrons traveling right to left

```

Fix the test

<< New golden standard

Create a test suite with these cases

Fancy cross

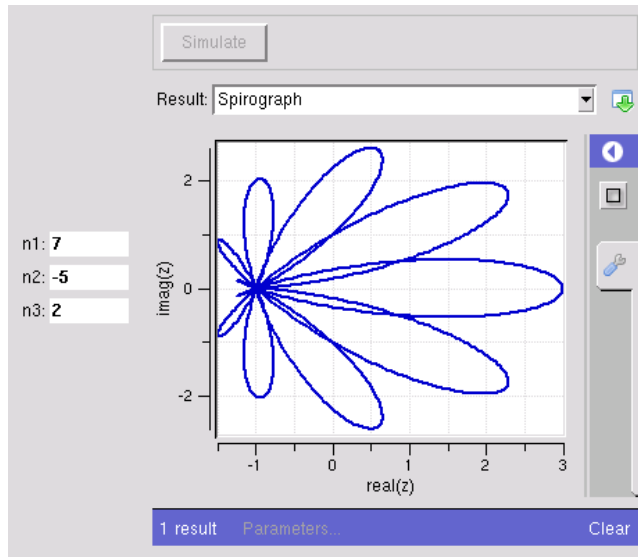
$$\begin{aligned} n_1 &= 13 \\ n_2 &= -7 \\ n_3 &= -3 \end{aligned}$$

Flower

$$\begin{aligned} n_1 &= 19 \\ n_2 &= -13 \\ n_3 &= 3 \end{aligned}$$

Palm Branch

$$\begin{aligned} n_1 &= 7 \\ n_2 &= -5 \\ n_3 &= 2 \end{aligned}$$



- Run the regression tests (should run cleanly)
- Edit one of the tests and delete some numbers
- Run the tests, discover the error, then regoldenize