Setting up PhysiCell on Windows



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@PhysiCell

PhysiCell Project

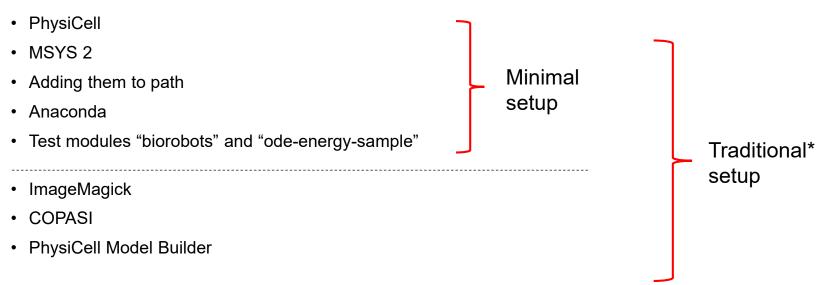
July 25, 2021



Thank you very much for watching!

Overview

This installation guide is prepared for installing PhysiCell to new fresh Windows. If you install PhysiCell 1.8.0 or previous version in your PC. Please uninstall MSYS and remove related directories from environmental variables.



^{*}Traditional setup includes optional tools that helps for pre-simulation edits and pro-simulation analysis

Youtube video

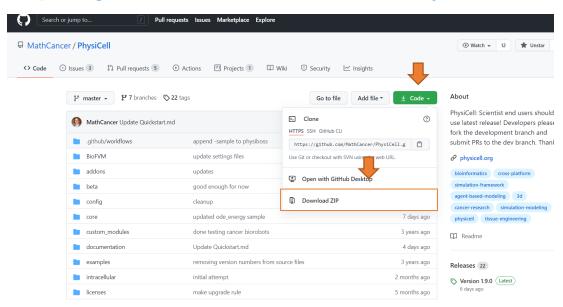
Here is the video link for the recording of installation on Windows.

LINK HERE!!!!!!!!!

PhysiCell

To download PhysiCell please proceed following link:

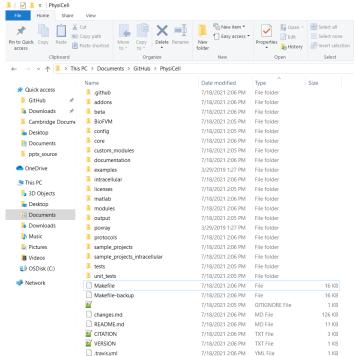
https://github.com/MathCancer/PhysiCell



- Please, click green "Code" button at right.
- Then, download as "ZIP".
- Extract to the desired directory.

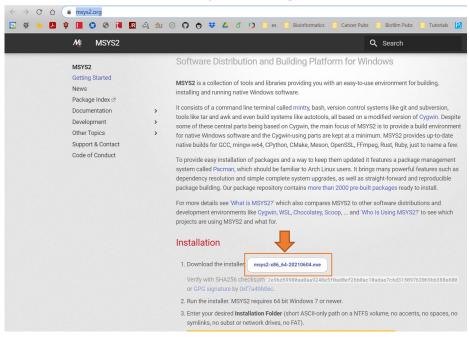
PhysiCell

After the extraction, PhysiCell folder should look like this:

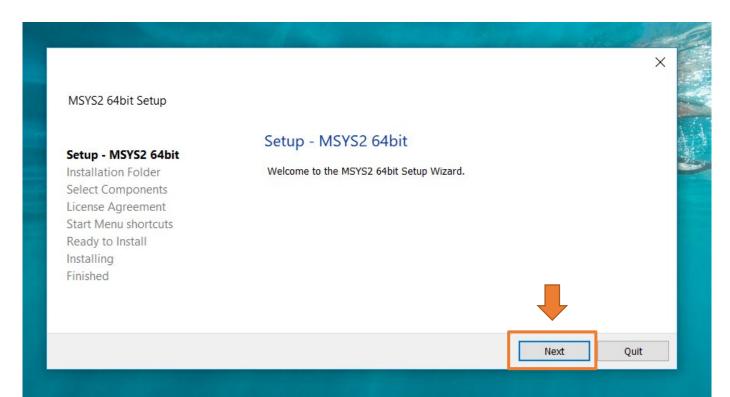


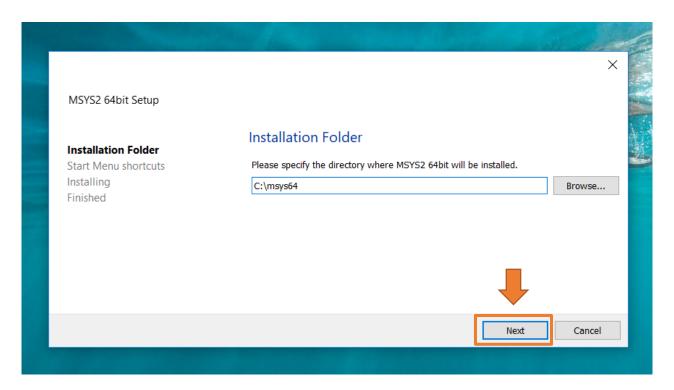
Please proceed following link:

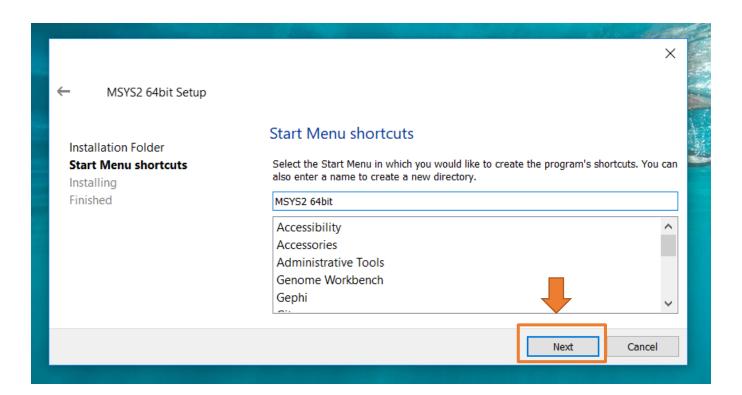
https://www.msys2.org/

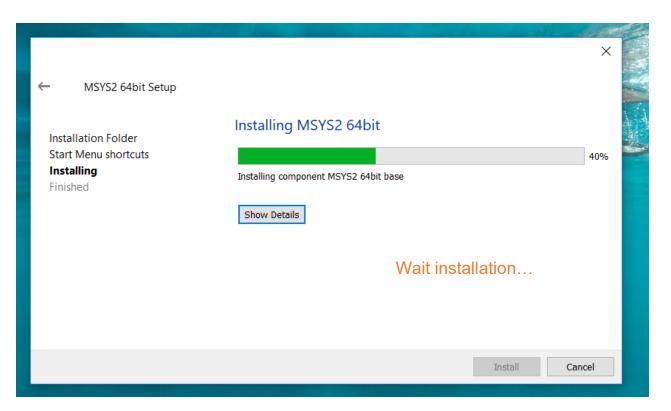


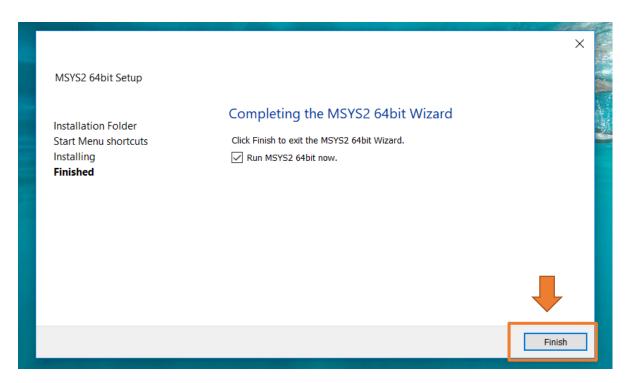
- Please, click "msys-x64_64-202XXX.exe" button at bottom.
- It should download installer.
- Open it with double-clicking the installer.





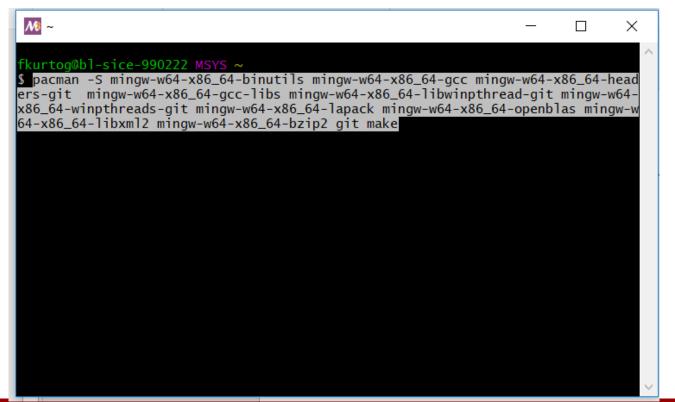






• After MSYS2 command prompt is opened, please copy and paste following line. Then, press enter.

```
pacman -S mingw-w64-x86_64-binutils mingw-w64-x86_64-gcc mingw-w64-x86_64-headers-git mingw-w64-x86_64-gcc-libs mingw-w64-x86_64-libwinpthread-git mingw-w64-x86_64-winpthreads-git mingw-w64-x86_64-lapack mingw-w64-x86_64-openblas mingw-w64-x86 64-libxml2 mingw-w64-x86 64-bzip2 git make
```



• Give "y" (yes) answer to the msys2. Press enter...

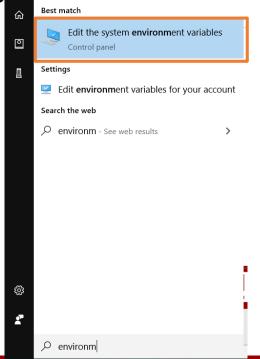
```
M₽ ~
                                                                              \times
             perl-HTTP-Cookies-6.10-1 perl-HTTP-Daemon-6.12-1
             perl-HTTP-Date-6.05-1 perl-HTTP-Message-6.32-1
             perl-HTTP-Negotiate-6.01-2 perl-IO-HTML-1.004-1
             perl-IO-Socket-SSL-2.070-1 perl-IO-Stringy-2.113-1
             perl-LWP-MediaTypes-6.04-1 perl-MIME-tools-5.509-1
             perl-MailTools-2.21-1 perl-Net-HTTP-6.21-1
             perl-Net-SMTP-SSL-1.04-1 perl-Net-SSLeay-1.90-1
             perl-TermReadKey-2.38-2 perl-TimeDate-2.33-1
             perl-Try-Tiny-0.30-1 perl-URI-5.09-1 perl-WWW-RobotRules-6.02-2
             perl-libwww-6.54-1 vim-8.2.2859-2 git-2.31.1-2 make-4.3-1
             mingw-w64-x86_64-binutils-2.36.1-3
             mingw-w64-x86_64-bzip2-1.0.8-2 mingw-w64-x86_64-gcc-10.3.0-2
             mingw-w64-x86_64-gcc-libs-10.3.0-2
             mingw-w64-x86_64-headers-git-9.0.0.6215.788f57701-1
             mingw-w64-x86_64-lapack-3.9.0-2
             mingw-w64-x86_64-libwinpthread-git-9.0.0.6215.788f57701-1
             mingw-w64-x86_64-libxml2-2.9.12-1
             mingw-w64-x86_64-openblas-0.3.13-2
             mingw-w64-x86_64-winpthreads-git-9.0.0.6215.788f57701-1
Total Download Size:
                       79.99 MiB
Total Installed Size: 645.91 MiB
: Proceed with installation? [Y/n] y
```

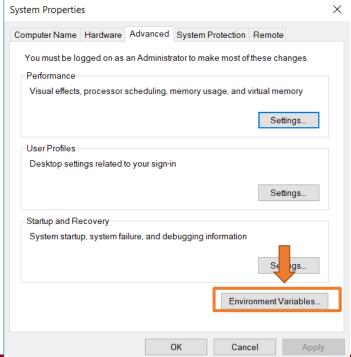
• It will take some time to install all required libraries. But in the end, you should not see any errors.

```
(45/57) installing perl-Try-Tiny
(46/57) installing perl-libwww
                                                  [################# 100%
Optional dependencies for perl-libwww
   perl-LWP-Protocol-https: for https:// url schemes
(47/57) installing perl-TimeDate
                                                   [################# 100%
(48/57) installing perl-MailTools
(49/57) installing perl-IO-Stringy
(50/57) installing perl-Convert-BinHex
                                                   [################### 100%
module test... pass.
(51/57) installing perl-MIME-tools
(52/57) installing perl-Net-SSLeay
(53/57) installing perl-IO-Socket-SSL
(54/57) installing perl-Net-SMTP-SSL
(55/57) installing perl-TermReadKey
                                                   ################### 100%
(56/57) installing git
                                                   `################## 100%
Optional dependencies for git
   python: various helper scripts
   subversion: git svn
(57/57) installing make
                                                  [########### 100%
:: Running post-transaction hooks...
(1/1) Updating the info directory file...
fkurtog@bl-sice-990222 MSYS ~
```

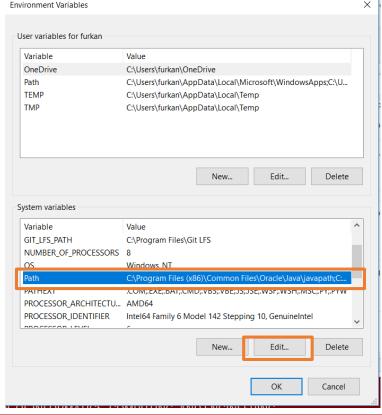
Please go to environmental variables with searching on Windows Start

Menu.





 Please select to SYSTEM PATH and press "Edit..."

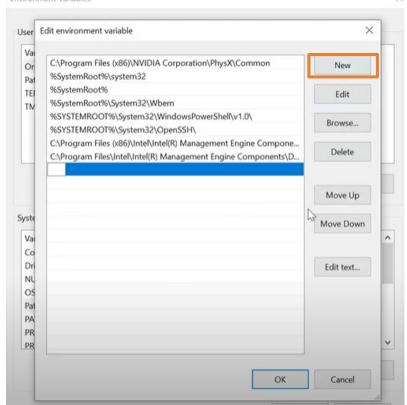


 Please press "New". And add following paths to there:

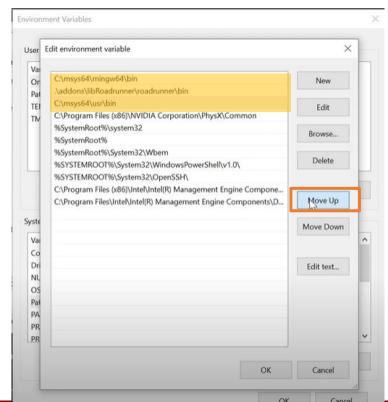
C:\msys64\mingw64\bin

C:\msys64\usr\bin

.\addons\libRoadrunner\roadrunner\bin



- Then please "MOVE UP", newly added three paths.
- The list should like the following image.



Please proceed following link:

https://www.anaconda.com/products/individual



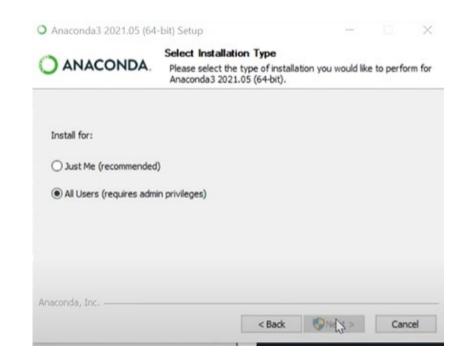
Individual Edition

Your data science toolkit

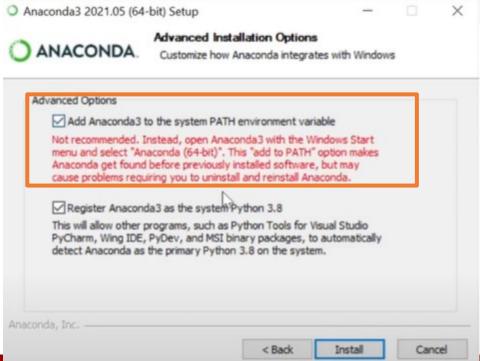
With over 25 million users worldwide, the open-source Individual Edition (Distribution) is the easiest way to perform Python/R data science and machine learning on a single machine. Developed for



- Please install for all users. (This is step is required for msys2 path priorities)
- If you cannot install for all users.
 Please add msys2 paths for
 current user in the environmental
 variables. (Instead of systems
 variables, please use upper user
 variables)



Please select option for adding Anaconda to the path.



Installation...

 Anaconda3 2021.05 (64-bit) Setup 			-		×
ANACONDA.	Installing Please wait while Anaconda3 2021.05 (64-bit) is being installed.				
Setting up the package car	he				
Show details					
Anaconda, Inc.		< Back	Next >	Cano	cel

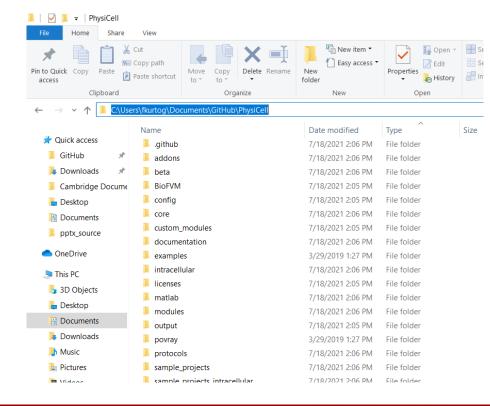
Test Module: "biorobots"

- At this moment PhysiCell is ready to work. Let's try "biorobots", first.
- Open command prompt from Start menu with typing "cmd".

```
rosoft Windows [Version 10.0.17134.1845]
 2018 Microsoft Corporation. All rights reserved
\Users\fkurtog>_
```

Going to PhysiCell folder

- Proceed to the PhysiCell folder at File Browser.
- Click to the directory address from top.
- · Copy it



Going to PhysiCell folder

• Type "cd" and paste the copied directory with space between them.

```
Command Prompt

Microsoft Windows [Version 10.0.17134.1845]

(c) 2018 Microsoft Corporation. All rights reserved.

C:\Users\fkurtog\cd C:\Users\fkurtog\Documents\GitHub\PhysiCell

C:\Users\fkurtog\Documents\GitHub\PhysiCell>_
```

Testing Modules (biorobots)

- Let's compile "biorobots" sample project first.
- Please run following commands in the command prompt.

```
make biorobots-sample
make
```

- This should successfully compiles biorobots sample project that ready to run with following command.
- ./biorobots.exe

First simulation...

```
Command Prompt - biorobots.exe
       Placing 15 director cells ...
       Placing cargo cells ...
       Placing worker cells ...
Using PhysiCell version 1.9.0
       Please cite DOI: 10.1371/journal.pcbi.1005991
       Project website: http://PhysiCell.MathCancer.org
See ALL_CITATIONS.txt for this list.
current simulated time: 0 min (max: 2880 min)
total agents: 514
interval wall time: 0 days, 0 hours, 0 minutes, and 0.0075989 seconds
total wall time: 0 days, 0 hours, 0 minutes, and 0.0094352 seconds
Using method diffusion_decay_solver__constant_coefficients_LOD_2D (2D LOD with Thomas Algorithm) ...
Warning and FYI: Very small vectors are normalized to 0 vector
current simulated time: 2 min (max: 2880 min)
total agents: 513
interval wall time: 0 days, 0 hours, 0 minutes, and 0.163112 seconds
total wall time: 0 days, 0 hours, 0 minutes, and 0.173366 seconds
current simulated time: 4 min (max: 2880 min)
total agents: 513
interval wall time: 0 days, 0 hours, 0 minutes, and 0.139892 seconds
total wall time: 0 days, 0 hours, 0 minutes, and 0.313664 seconds
```

Testing Modules (ode-energy-sample)

• If the first test is successfully created .mat and .xml files in the output folder. Let's try "ode-sample" sample project. To do that:

```
make ode-energy-sample
make
```

• While compilation, you should see "libroadrunner" installation. After the compilation, please use following command to run simulation.

```
./ode energy.exe
```

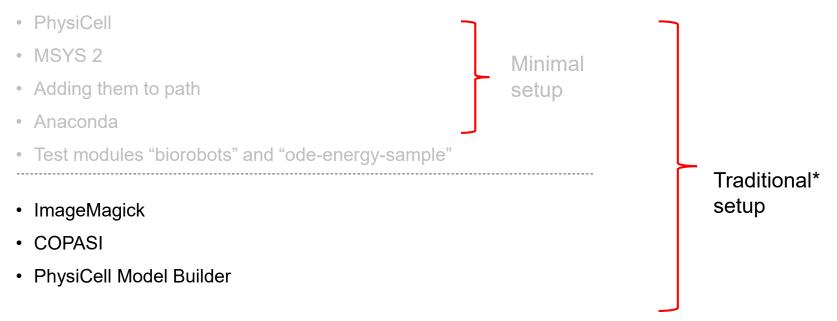
Second simulation... (ode-energy-sample)

```
    Command Prompt - ode_energy.exe

Number of boundary species = 0
lumber of compartments = 1
loating species names:
Glucose Oxygen Energy Lactate apoptosis_rate migration_speed Lac_Secretion_Rate Transition_Rate
Glucose -> 0
Oxygen -> 1
nergy -> 2
actate -> 3
poptosis rate -> 4
nigration_speed -> 5
_ac_Secretion_Rate -> 6
Transition Rate -> 7
100
  0.000166667
----- start: librr_intracellular.cpp: start() called
 ------ start: doing: rrHandle = createRRInstance()
 ----- start: rrHandle = 0x1d1bb045f30
    sbml_filename = ./config/Toy_Metabolic_Model.xml
```

Overview

We have finished the minimal setup. Following slides will show traditional setup.



*Traditional setup includes optional tools that helps for pre-simulation edits and pro-simulation analysis



ImageMagick

- Please proceed the following link: https://imagemagick.org/script/download.php
- And go below for "Windows Binary Release" and click there to download.

Windows Binary Release

ImageMagick runs on Windows 10 (x86 & x64), Windows 8 (x86 & x64), Windows 7 (x86 & x64), Windows Server 2012, Windows Vista (x86 & x64) with Service Pack 2, Windows Server 2008 (x86 & x64) with Service Pack 2, and Windows Server 2008 R2 (x64).

The amount of memory can be an important factor, especially if you intend to work on large images. A minimum of 512 MB of RAM is recommended, but the more RAM the better. Although ImageMagick runs well on a single core computer, it automagically runs in parallel on multi-core systems reducing run times considerably.

The Windows version of ImageMagick is self-installing. Simply click on the appropriate version below and it will launch itself and ask you a few installation questions. Versions with *Q8* in the name are 8 bits-per-pixel component (e.g. 8-bit red, 8-bit green, etc.), whereas, *Q16* in the filename are 16 bits-per-pixel component. A Q16 version permits you to read or write 16-bit images without losing precision but requires twice as much resources as the Q8 version. Versions with *dll* in the filename include ImageMagick libraries as <u>dynamic link libraries</u>. Unless you have a Windows 32-bit OS, we recommend this version of ImageMagick for 64-bit Windows:

Version	Description
ImageMagick-7.1.0-4-Q16-HDRI-x64-dll.exe	Win64 dynamic at 16 bits-per-pixel component with High-dynamic-range imaging enabled

Or choose from these alternate Windows binary distributions:





ImageMagick

Please be sure the option for adding directory to path is selected.

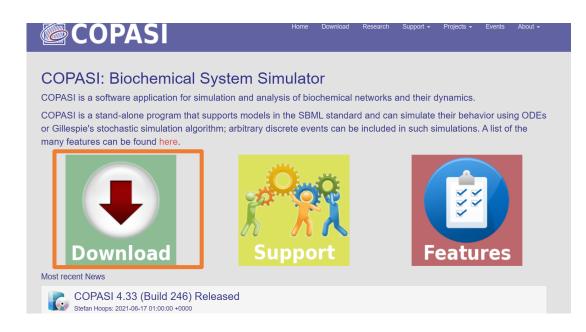
Select Additional Tasks			-
Which additional tasks should be performed	ed?		(HOI)
Select the additional tasks you would like 7.1.0 Q16-HDRI (64-bit), then click Next.	Setup to perform w	hile installing I	mageMagick
Create a desktop icon			
Add application directory to your system	em path		
Install rempeg			
Associate supported file extensions wi	th ImageMagick		
Install legacy utilities (e.g. convert)			
☐ Install development headers and librar	ries for C and C++		
☐ Install PerlMagick for Strawberry Perl	v5.20		
Install ImageMagickObject OLE Contro	ol for VBscript, Visu	al Basic, and W	SH
	Back	Next.	Cancel

COPASI

Please proceed following link to download COPASI

http://copasi.org/

Install regularly



PhysiCell Model Builder (1)

- The Model Builder is a GUI to let you create/edit a .xml configuration file that defines (nearly all of) a PhysiCell model.
- Download the latest release at:

https://github.com/PhysiCell-Tools/PhysiCell-model-builder/releases

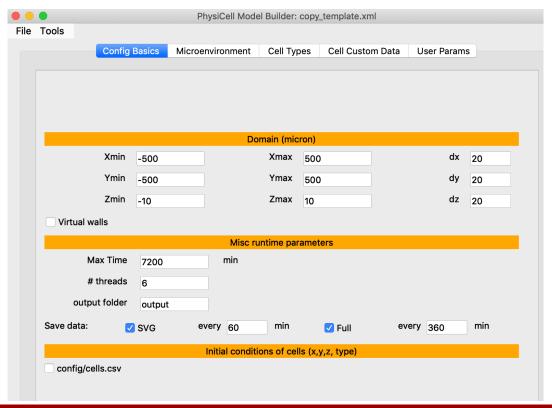
Uncompress the .zip, change directory into it, and run it:

```
$ unzip PhysiCell-model-builder-1.1.zip
$ cd PhysiCell-model-builder-1.1
$ python bin/gui4xml.py
```

This should display the GUI (next page):

Note that if your setup automatically unzips files after downloading, this command will not work. INSTEAD, **skip** the unzip command and go directly to the second command

PhysiCell Model Builder (2)



A User Guide for the Model Builder is still be written.

It is a tool that is still considered "beta", so your feedback will be very valuable.

Funding Acknowledgements







PhysiCell Development:

- Breast Cancer Research Foundation
- Jayne Koskinas Ted Giovanis Foundation for Health and Policy
- National Cancer Institute (U01CA232137)
- National Science Foundation (1720625)

Training Materials:

Administrative supplement to NCI U01CA232137 (Year 2)