



Reaction Simulator:

A BIOEN 437 project

By: Matthew Van Ginneken & Olivia Walsh

Background

- Create a web based app to visualize reaction kinetics
 - Easy to use for beginners
 - Accessible
 - Simple
- Similar Packages
 - COPASI
 - JigCell

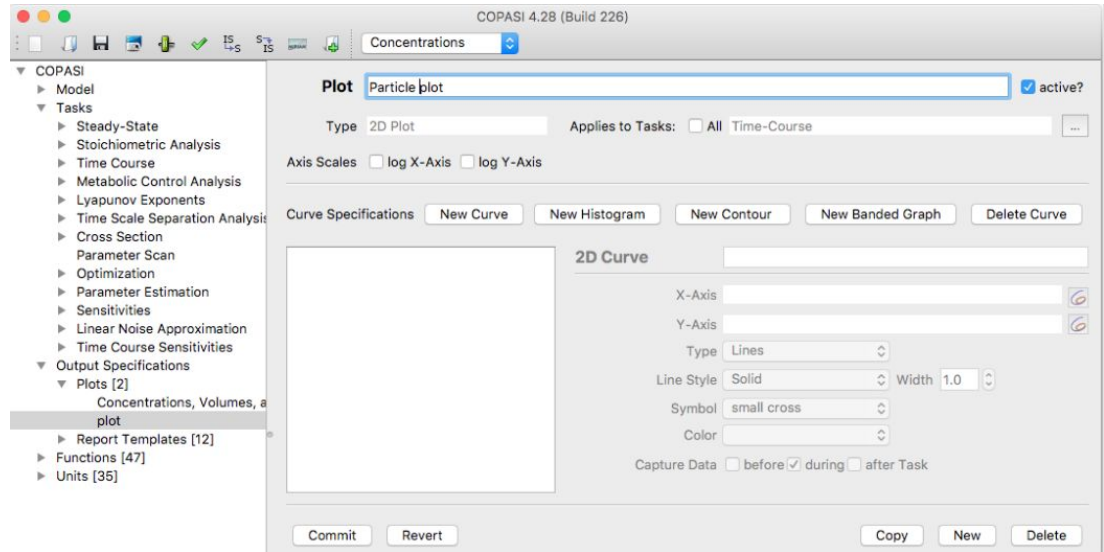


Figure 1: A clip of COPASI graph creator



Use Cases

- 1) The user can iteratively submit reactions and constants and then view the steady state graph
- 2) The user can submit an antimony txt file and then view the steady state graph and steady state values
- 3) The user can submit an SBML file and then view the steady state graph and steady state values

Demo

Welcome to the Tellurium Model Builder. Here you can input reactions one at a time in order to build a Computational Biology Mass Action Model.

Add a reaction to your model

Reactant and Product names should be entered as a comma separated list of strings. i.e : A,B,C

If none of the Reactant or Products are fixed enter "None" into the field

Starting Concentrations should be entered as a comma separated list of floats. i.e : 0.45,1.34,5

The reaction constant should be entered as a single float unless the reaction is reversible in which case

enter the reaction constants as 2 floats separated by a comma i.e : 0.23 or 0.23,0.12

If your reaction is reversible check the Reversible box

Reactant Names:

Starting Reactant Concentrations:

Fixed Reactant Names:

Product Names:

Starting Product Concentrations:

Fixed Product Names:

Reaction Constant:

Reversible ☐

Press "Add Reaction" when you have finished inputting your data for the reaction

Add Reaction

Press "Run Simulation" when you have added all your reactions to the model

Run Simulation

If you need to reset the model press "Reset Model" which will delete all reactions currently stored in the model

Reset Model

Upload an Antimony Text File Bars.txt

This field is required.

SBML File Link

User Runs Python Files Design - Flask Webpage

python initializes Flask App
And hosts it on the local web
port.

User navigates to web port
`http://127.0.0.1:5000`

System Rests
Directs user to Builder



Reset Model

Add Reaction

Model Upload

Submit SBML File

Run Simulation

uses Python Func() to reset model

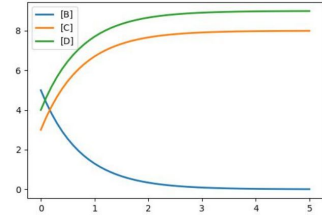
Sends user input to Save2Dict

Runs loadModel()

Runs loadSBML()

Runs runSim()

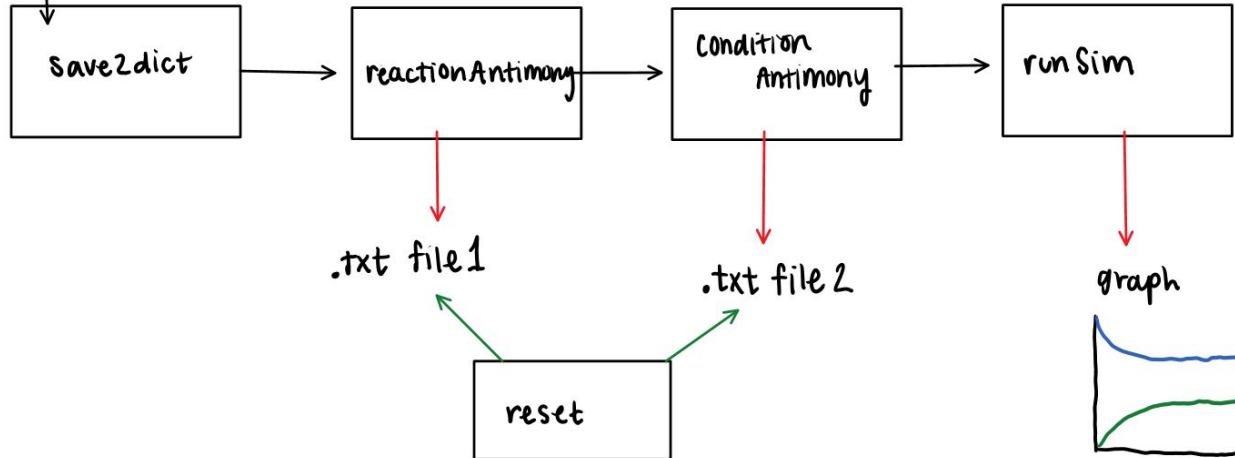
Steady State Plot Image



Simulation is Run and
SS Graph is Produced

Design - Running the Simulations Through Python

Flask webpage
user inputs
strings





Design - Python Functions

- Save2dict
- Antimony Builder
 - 2 separate functions, iteratively add to two separate txt files

```
2  $A+B+C->E ; k0*A*B*C
3  E->$A+B+C ; k1*E
```

```
2  A=10;B=0;C=0
3  E=3
4  k0=10;k1=2
```

```
2  dict = {
3      'Reactants' = ['A', 'B'],
4      'fixedReactants' = ['A'],
5      'ReactantIC' = [10,2],
6      'Products' = ['C'],
7      'fixedProducts' = ['C'],
8      'ProductIC' = [0],
9      'Reversibility' = False,
10     'RxnConstant' = [3]
```

- runSim
 - JPG file of the graph

Project Structure

AMCOWMVG.egg-info	Working Pip Installable Version
AMCOWMVG	Working Pip Installable Version
__pycache__	Testing2
dist	Working Pip Installable Version
docs	Final Final Version
examples	Working Pip Installable Version
LICENSE	Testing2
MANIFEST.in	Working Pip Installable Version
README.md	Final Final Version
pyproject.toml	Working Pip Installable Version
setup.cfg	Working Pip Installable Version
test_unit.py	Testing2

```
D:.\
|  LICENSE
|  MANIFEST.in
|  pyproject.toml
|  README.md
|  setup.cfg
|  test_unit.py
|
+---AMCOWMVG
|  |  antimonyTools.py
|  |  forms.py
|  |  routes.py
|  |  testing.py
|  |  __init__.py
|  |
|  +---static
|  |  |  antimony1.txt
|  |  |  antimony2.txt
|  |  |  output.jpg
|  |  |  test.txt
|  |
|  +---templates
|  |  |  buildSystem.html
|  |  |  home.html
|  |  |  results.html
|  |  |  testing.html
|  |
+---docs|
|  |  Component Specifications.pdf
|  |  Final Presentation.pdf
|  |  Functional Specs_updated.pdf
|  |
+---examples
|  |  Antimony_Text_Upload.mp4
|  |  Base_Case.mp4
|  |  reset.mp4
|  |  SBML_Upload.mp4
```




Lessons Learned

- Working on different parts of the code then combining
- Creating tests using Python's *unittest*
- Documentation and process of creating working code and submitting as a python package



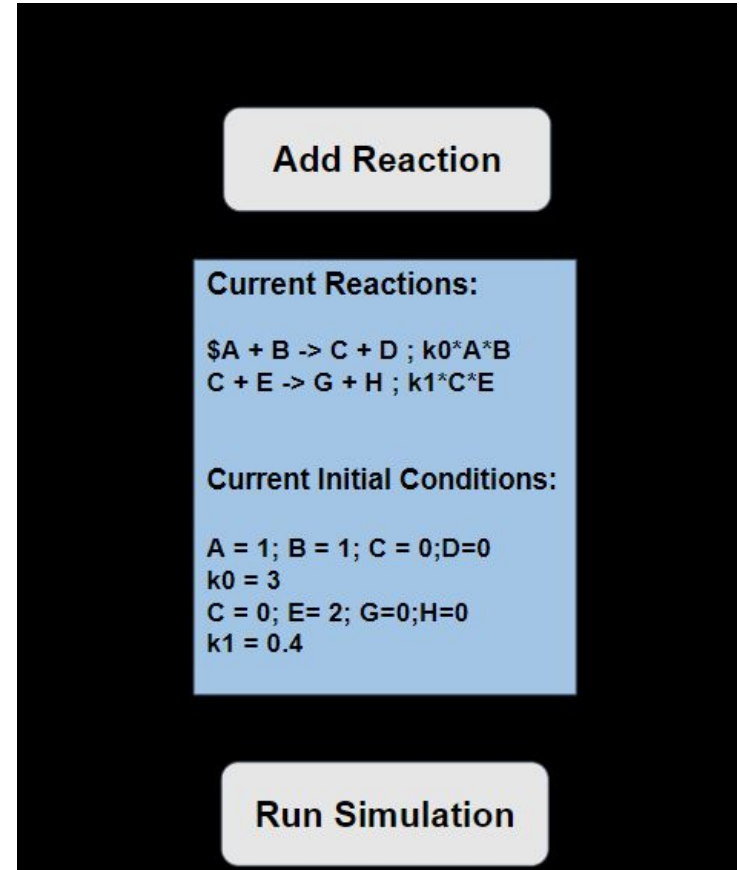
Future Directions

- Adding in other functions from tellurium
 - Showing different features of the simulation such as steady state values and system fluxes
- User can export their antimony txt file and then resume working with it on our webpage
- Being able to remove a reaction after it has been submitted

Future Directions

Listing the reactions that are submitted

- Help the user with organization
- User can check if the reaction was inputted correctly





Future Directions

Allowing the user to simulate more than one type of reaction kinetics

- Implemented with options given in a home page

What type of kinetics would you like to model?

Michaelis Menten Kinetics

Mass Action Kinetics

Brigg – Haldane Steady State (Enzyme)

Rapid Equilibrium Approximation (Enzyme)

Reversible Enzyme Catalysis