https://github.com/physicell-training/ws2023

# Lesson 7: Working with PhysiCell Data in Python



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# **PhysiCell Project**

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# pcdl - "PhysiCell's USB port"

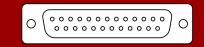
### Focus:

- stay backwards compatible, as much as possible.
- stay platform independent (Windows, MacOS, Linux).
- be **pip** installable.
- load PhysiCell output into Python3.
- supply the loaded data in well established, standardized data formats.
  - pandas DataFrame and Series
  - scverse **AnnData** objects

#### In other words:

- pcdl does not try to be analysis software.
- pcdl tries to be the default connector between PhysiCell output and analysis software.

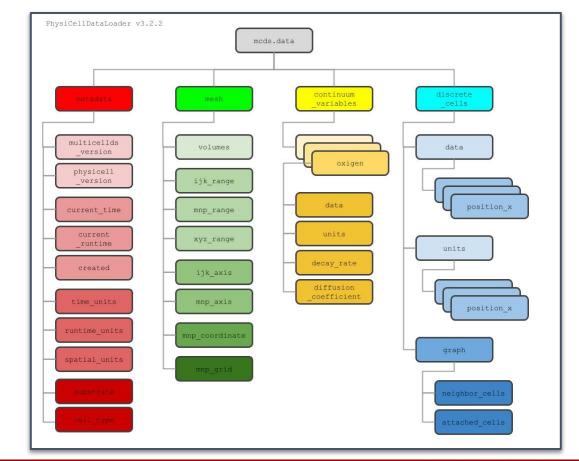
In this presentation, we will see, how far we come with this strategy.



# **Blueprint**

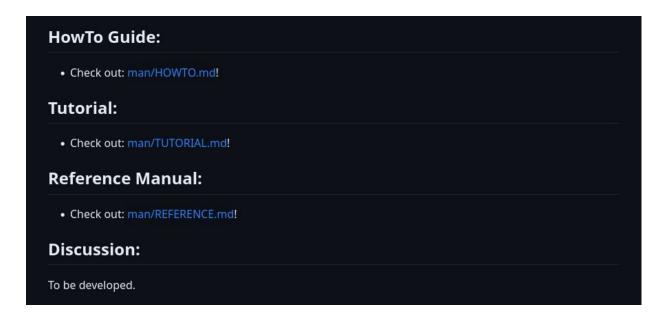
one time step =
one mcds object =
one Python3 dictionary
and a set of functions to
access the data.

mcds: Multi Cellular Data Standard



# pcdl - all the details

https://github.com/elmbeech/physicelldataloader/blob/master/README.md



# pcdl[all] - python library dependencies

### pip install pcdl # light weight, the basics:

- matplotlib
- numpy
- pandas
- scipy

### pip install pcdl[scverse] # heavy weight:

anndata - for single cell data analysis

### pip install pcdl[data] # used for the tutorial:

requests - for http downloading the data files



# pcdl - the workhorse functions for data analysis

```
TimeStep Class
 help(pcdl.TimeStep) # ! make class instance
 help(pcdl.TimeStep. init )
 # TimeStep microenvironment
 help(pcdl.TimeStep.get concentration df) # ! workhorse function
 help(pcdl.TimeStep.get_conc_df) # ! shorthand
 help(pcdl.TimeStep.get_contour) # ! workhorse function
 # TimeStep cells and other agents
 help(pcdl.TimeStep.get cell df) # ! workhorse function
 help(pcdl.TimeStep.get_scatter) # ! workhorse function
 # TimeStep unit
 help(pcdl.TimeStep.get_unit_se) # ! workhorse function
 # TimeStep anndata
 help(pcdl.TimeStep.get anndata) # ! workhorse function
```

# pcdl - the workhorse functions for data analysis

```
TimeSeries Class
                                                                                                                   Q
 help(pcdl.TimeSeries) # ! make class instance
 help(pcdl.TimeSeries._init_)
 # TimeSeries load data
 help(pcdl.TimeSeries.get_mcds_list) # ! workhorse function
 help(pcdl.TimeSeries.get_anmcds_list) # ! workhorse function
 # TimeSeries triage data
help(pcdl.TimeSeries.get_cell_df_states) # ! workhorse function
 help(pcdl.TimeSeries.get_conc_df_states) # ! workhorse function
 # TimeSeries images and movies
 help(pcdl.TimeSeries.make imgcell) # ! workhorse function
help(pcdl.TimeSeries.make_imgsubs) # ! workhorse function
 help(pcdl.TimeSeries.make_gif) # ! workhorse function
 help(pcdl.TimeSeries.make_movie) # ! workhorse function
 # TimeSeries anndata
help(pcdl.TimeSeries.get_anndata) # ! workhorse function
```



# pcdl - data analysis...

- is not **source code** programming! (Although, when you write an analysis tool, you should use pcdl to load PhysiCell output.)
- is interactive "REPL (read-eval-print loop) programming".
   REPL programming is one-liner coding.
   REPL programming is ideal for ad hoc, on the fly data analysis.
   REPL programming is more agile than any GUI (graphical user interface) can be.

In the beginning, I will move slowly, to show how I do REPL programming. Afterward I will move fast, to give an overview about what is possible.

# live demo! pcdl\_ws2023.ipynb

https://github.com/elmbeech/physicelldataloader/blob/master/man/jupyter/pcdl\_ws2023.ipynb

### Conclusion

pcdl is simply and interface.

 don't re-invent the wheel: it is a fruitful approach to learn, make use of, and contribute to in the filed well established high-level Python3 libraries.

# Where to go from here?

#### the basics:

- learn core python3: https://www.python.org/ [resources: https://nostarch.com/python-kids-2nd-edition, https://realpython.com/]

- learn numpy: https://numpy.org/ learn scipy: https://scipy.org/ learn pandas: https://pandas.pydata.org learn matplotlib: https://matplotlib.org/

#### depending on your needs, learn:

- statsmodels: https://www.statsmodels.org/stable/index.html
- sklearn: https://scikit-learn.org/stable/
- pytorch: <a href="https://pytorch.org/">https://pytorch.org/</a> skimage: <a href="https://scikit-image.org/">https://scikit-image.org/</a>

#### at https://scverse.org:

- read the anndata docs the basic data object read the scanpy docs single cell analysis read the squidpy docs spatial single cell analysis
- read the scvi-tools docs single cell machine learning

#### at https://github.com/PhysiCell-Tools/python-loader:

- work through the pcdl TUTORIAL.md
- read the docstrings from the "workhorse functions" mentioned in the pcdl REFERENCE.md



# pcdl - future development

### **Road Map:**

- · vtk file format output, maybe stl and wavefront obj output.
- GML (networkx and igraph compatible) output.

# Acknowledgement

The whole **MathCancer** lab!

### python-loader original implementation:

- Pat Wall
- Randy Heiland Paul Macklin

### pcdl 2022 / 2023 evolution:

- Marshal Gress
- Heber Rocha

### OHSU:

- Jenny Eng (scanpy)
- Tina Ghodsi Asnaashari (abm)

**Software** - because pcdl is standing on the shoulder of giants:

- PhysiCell,
- PhysiCell Studio
- Python3 core libraries
- numpy, scipy, pandas, matplotlib
- scveres: anndata
- http: requests

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