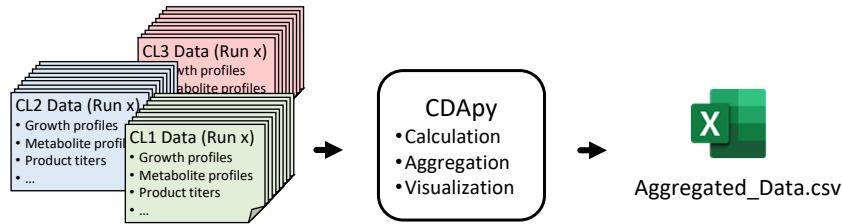


Cell Culture Data Pipeline Python Package (CDAPy)



Process Data Entry (Excel)

Multiple batches and cell lines

New features compared to Excel worksheet:

- Process multiple batches and cell lines at the same time
- Provide rolling window regression for specific-rate calculation
- Provide interactive plotting to analyze and compare process profiles

Credit to Yudai Fukae

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Excel data input (1): process data entry

Data (Date/Run/Cell Line)	Cell Line	Experiment Data					Feed Volume (mL)			Concentration Before Feeding				Concentration After Feeding			
		ID	Initial Volume (mL)	Sample Volume (mL)	Time Added (min)	Viability Cell Concentration (10 ⁶ cells/mL)	Dwell Cell Concentration (10 ⁶ cells/mL)	Total Cell Concentration (10 ⁶ cells/mL)	AGI (mg/L)	F01	F02	Feed 3	Aspartate (mM)	Glucose (mM)	Glutamine (mM)	Lactate (mM)	Lactate (mM)
CL1	74.00	1.00	0.00	0.1870	0.0000	0.1870	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	0.4030	0.0000	0.4030	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	0.8970	0.0000	0.8970	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	1.6200	0.0014	1.6214	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	3.2600	0.0096	3.2696	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	5.1200	0.0265	5.1265	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	5.1400	0.0407	5.1807	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	17.4000	0.0587	17.4587	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	18.5000	0.0593	18.5593	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	20.5000	0.0593	20.5593	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	18.0000	0.1130	18.1130	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	18.4000	0.0558	18.4558	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	15.4000	0.2354	15.6354	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	0.4847	0.0007	0.4854	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	0.6769	0.0007	0.6776	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	1.0469	0.0015	1.0484	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	1.4953	0.0051	1.5004	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	2.1107	0.0184	2.1291	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	3.5471	0.0204	3.5675	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	3.4855	0.0195	3.5050	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	3.8142	0.0980	3.9122	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	3.9834	0.1262	4.1096	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	4.0619	0.1367	4.1986	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	4.1798	0.1284	4.3082	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	4.0797	0.1698	4.2495	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
CL1	74.00	1.50	0.00	4.0046	0.1469	4.1515	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000

- Enter basic process data (orange, excluding concentrations), feed volume (green), concentration before feeding (blue), and concentration after feeding (yellow, optional)
- Accept datasets from cell lines/processes that have different settings (e.g., different feeds or measurements)
- Can add columns to include more online/offline measurements (orange), feeds (green), or chemical species (blue and orange)

* This Excel template is different from the previous Excel batch sheet.

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Excel data input (2): feed concentration and polynomial degree sheets

	A	F	G	I
1	Cell Line	CL1	CL1	CL2
2	ID	13	13	21
3	Feed Name	F01	F02	Feed 2
9	Glucose (mM)	400.0		400.0
10	Glutamine (mM)		200.0	
11	Glutamate (mM)	11.6		
15	Lactate (mM)	0.0		

- Enter the composition of feeds in the “Feed Concentration” sheet
- Feed name should be consistent with the ones in the “Measured Data” sheet

	A	D	E
1	Cell Line	CL1	CL2
2	ID	13	21
3	Cell	3	3
9	Glucose	4	4
10	Glutamine	3	3
11	Glutamate	4	4
15	Lactate	5	6

- Define the polynomial degree used for the polynomial regression of specific rate calculations
- Can use varying degrees for different species and cell lines/experiments (IDs)

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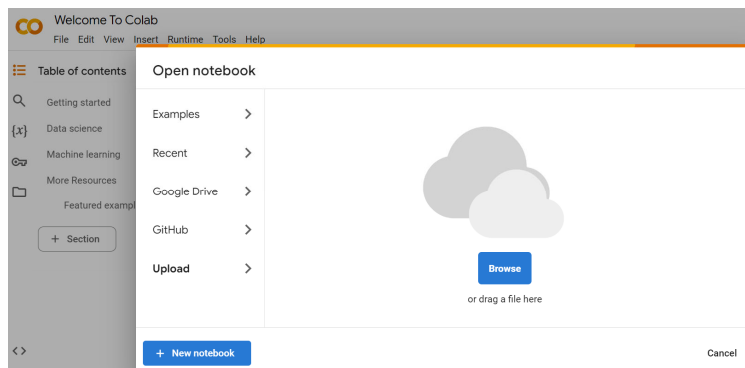
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Step-by-step to run tutorial notebooks on Google Colab (1)

Step 1: Open Google Colab (<https://colab.research.google.com/>)

* Google Colab is a hosted Jupyter Notebook service that requires no setup to use and provides free-of-charge access to computing resources

Step 2: Upload the “Part2_Python_oackage_tutorial.ipynb” to Google Colab



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Step-by-step to run tutorial notebooks on Google Colab (2)

Step 3: Run the first cell (grey area shown below) with the “play” bottom to setup the package

1. Package setup

💡 Can ignore the printout when executing the code unless error message raises.

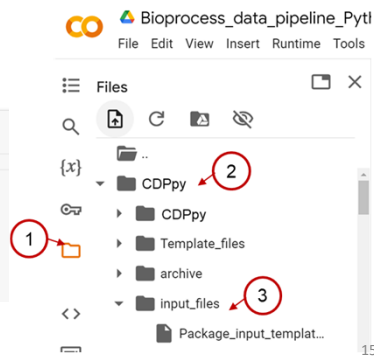
Play

```
# clone package from GitHub
!git clone https://github.com/ddolab/CDPpy.git

# install/update necessary package for plotting
!pip install dash
!pip install pandas==1.4.3

# go into the package folder
import os
os.chdir('/content/CDPpy')
```

Step 4: Follow the steps in the screenshot and upload the input Excel file into the “input_files” folder



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Step-by-step to run tutorial notebooks on Google Colab (3)

Step 5: type the file name of the input Excel file in section 2.1

```
# setup the input filename
input_fname = 'Part2_Package_input_Example.xlsx'
```

Step 6: Performing the data processing by running the rest of the notebook with “play” bottoms **Play**

- Section 2: example of processing one cell line data
- Section 3: example of processing multiple cell line data (slide 17)

Step 7: Go to section 4 to plot and analyze process profiles. (Slide 18)

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Full codes for processing multi-cell-line datasets in CDAPy

```
# Specify parameters that store data information for cell line 1 (CL1)
param_1 = FedBatchParameters(
    # input the cell line name, be consistent with the input excel file
    cell_line_name='CL1',
    # input "True" if there are measurements on concentrations after feeding, otherwise, "False"
    use_concentration_after_feed=False,
    # input "True" if feeding composition is known, otherwise, "False"
    use_feed_composition=True)

# Specify parameters that store data information for cell line 2 (CL2)
param_2 = FedBatchParameters(
    # input the cell line name, be consistent with the input excel file
    cell_line_name='CL2',
    # input "True" if there are measurements on concentrations after feeding, otherwise, "False"
    use_concentration_after_feed=False,
    # input "True" if feeding composition is known, otherwise, "False"
    use_feed_composition=True)
```

[Specify data information for each cell line](#)

```
### Repeat section 2.2
# create a Python object to preform the data processing of CL1/2
cell_line_1_2 = FedBatchCellCulture() **Change input filename if needed

# define path to the input data file. Change the filename if needed.
path = input_path(input_fname)

# load the data set to the data-processing object
cell_line_1_2.load_data(file=path)

# perform data processing with specified parameters that store data information.
# pass different data information with separate input params
cell_line_1_2.perform_data_process(parameters=[param_1, param_2])
```

[Input and process data](#)

```
# export the processed dataset
cell_line_1_2.save_excel(file_name='output_CL1_2.xlsx')
```

[Save results to Excel](#)

* More details can be found in sections 2 & 3 in tutorial Jupyter notebook

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Interactive Plotting Widget – Customized Data Visualization

[Select data to be visualized](#)

Cell Line: Experiment ID:

Profile:

Species: SP. Rate Method:

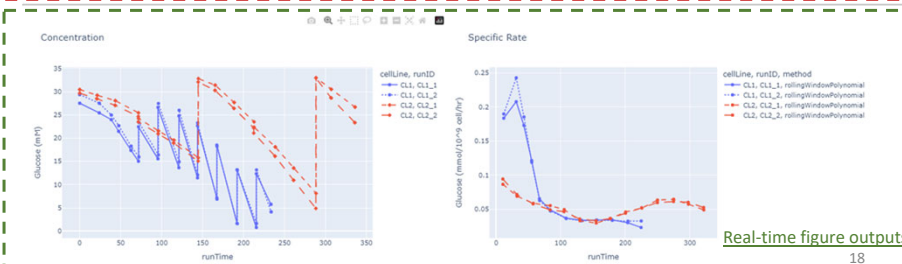
Concentration Graph Style: ☐ Cell Line ☐ Run ID Color: ☐ Cell Line ☐ Run ID Line: ☐ Cell Line ☐ Run ID Symbol: ☐ Cell Line ☐ Run ID

Cumulative Concentration Graph Style: ☐ Cell Line ☐ Run ID Color: ☐ Cell Line ☐ Run ID Line: ☐ Cell Line ☐ Run ID Symbol: ☐ Cell Line ☐ Run ID

Specific Rate Graph Style: ☐ Cell Line ☐ Run ID ☐ Method Color: ☐ Cell Line ☐ Run ID ☐ Method Line: ☐ Cell Line ☐ Run ID ☐ Method Symbol: ☐ Cell Line ☐ Run ID ☐ Method

Legend: ☐ on ☐ off

[Customize figure styles](#)



[Real-time figure outputs](#)

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