

import sys

BPL_TEST2_Batch_calibration script with PyFMI

The key library PyFMI is installed.

After the installation a small application BPL_TEST2_Batch_calibration is loaded and run. You can continue with this example if you like.

```
In [1]:
        !lsb_release -a # Actual VM Ubuntu version used by Google
       No LSB modules are available.
       Distributor ID: Ubuntu
                      Ubuntu 22.04.4 LTS
       Description:
       Release:
                       22.04
       Codename:
                       jammy
In [2]: %env PYTHONPATH=
       env: PYTHONPATH=
        !python --version
In [3]:
       Python 3.11.11
In [4]: !wget https://repo.anaconda.com/miniconda/Miniconda3-py311_24.11.1-0-Linux-x86_64.s
        !chmod +x Miniconda3-py311_24.11.1-0-Linux-x86_64.sh
        !bash ./Miniconda3-py311_24.11.1-0-Linux-x86_64.sh -b -f -p /usr/local
```

sys.path.append('/usr/local/lib/python3.11/site-packages/')

```
--2025-03-26 10:26:56-- https://repo.anaconda.com/miniconda/Miniconda3-py311_24.11.
       1-0-Linux-x86_64.sh
       Resolving repo.anaconda.com (repo.anaconda.com)... 104.16.32.241, 104.16.191.158, 26
       06:4700::6810:bf9e, ...
       Connecting to repo.anaconda.com (repo.anaconda.com) | 104.16.32.241 | :443... connected.
       HTTP request sent, awaiting response... 200 OK
       Length: 145900576 (139M) [application/octet-stream]
       Saving to: 'Miniconda3-py311_24.11.1-0-Linux-x86_64.sh'
       Miniconda3-py311_24 100%[==========>] 139.14M 144MB/s
                                                                          in 1.0s
       2025-03-26 10:26:57 (144 MB/s) - 'Miniconda3-py311_24.11.1-0-Linux-x86_64.sh' saved
       [145900576/145900576]
       PREFIX=/usr/local
       Unpacking payload ...
       Installing base environment...
       Preparing transaction: ...working... done
       Executing transaction: ...working... done
       installation finished.
In [5]: !conda update -n base -c defaults conda --yes
```

Channels:

- defaults

Platform: linux-64

Collecting package metadata (repodata.json): - 22\ 22| 22/ 22- 22\ 22| 22/ 22- 22\

22 | 22 / 22 - 22 \ 22 | 22 / 22done Solving environment: \ 22 | 22done

Package Plan

environment location: /usr/local

added / updated specs:

- conda

The following packages will be downloaded:

package	build	
ca-certificates-2025.2.25 certifi-2025.1.31 openssl-3.0.16	h06a4308_0 py311h06a4308_0 h5eee18b_0	129 KB 163 KB 5.2 MB
	Total:	5.5 MB

The following packages will be UPDATED:

Downloading and Extracting Packages:

```
openssl-3.0.16 | 5.2 MB | : 0% 0/1 [00:00<?, ?it/s] certifi-2025.1.31 | 163 KB | : 0% 0/1 [00:00<?, ?it/s] ca-certificates-2025 | 129 KB | : 0% 0/1 [00:00<?, ?it/s] ca-certificates-2025 | 129 KB | : 100% 1.0/1 [00:00<00:00, 25.64it/s] certifi-2025.1.31 | 163 KB | : 100% 1.0/1 [00:00<00:00, 19.60it/s]
```

ca-certificates-2025 | 129 KB | : 100% 1.0/1 [00:00<00:00, 12.44it/s]

```
Preparing transaction: - 22done
Verifying transaction: | 22/22-22done
Executing transaction: | 22done
```

```
In [6]: !conda --version
!python --version
```

```
conda 24.11.1
Python 3.11.11
```

In [7]: !conda config --set channel_priority strict

In [8]: !conda install -c conda-forge pyfmi --yes # Install the key package

Channels:

- conda-forge

defaults

Platform: linux-64

Collecting package metadata (repodata.json): - 22\ 22| 22/ 22- 22\ 22| 22/ 22- 22\ 22| 22/ 22- 22\ 22| 22/ 22- 22\ 22| 22/ 22- 22\ 22|

⊡done

Solving environment: - ₽₽\ ₽₽| ₽₽done

Package Plan

environment location: /usr/local

added / updated specs:

- pyfmi

The following packages will be downloaded:

package	build			
_x86_64-microarch-level-3	2_broadwell	8	KB	conda-forge
assimulo-3.6.0	py311h083bc19_0	1.1	MB	conda-forge
certifi-2025.1.31	pyhd8ed1ab_0	159	KB	conda-forge
conda-25.1.1	py311h38be061_1	1.1	MB	conda-forge
fmilib-2.4.1	hac33072_1	383	ΚB	conda-forge
gmp-6.3.0	hac33072_2	449	KB	conda-forge
libamd-3.3.3	haaf9dc3_7100102	49	KB	conda-forge
libblas-3.9.0	31_h59b9bed_openblas		16	KB conda-forge
libbtf-2.3.2	h32481e8_7100102	27	KB	conda-forge
libcamd-3.3.3	h32481e8_7100102	46	ΚB	conda-forge
libcblas-3.9.0	31_he106b2a_openblas		16	KB conda-forge
libccolamd-3.3.4	h32481e8_7100102	42	ΚB	conda-forge
libcholmod-5.3.1	h59ddab4_7100102	1.1	MB	conda-forge
libcolamd-3.3.4	h32481e8_7100102	33	ΚB	conda-forge
libcxsparse-4.4.1	h32481e8_7100102	118	ΚB	conda-forge
libgcc-14.2.0	h767d61c_2	828	ΚB	conda-forge
libgcc-ng-14.2.0	h69a702a_2	52	KB	conda-forge
libgfortran-14.2.0	h69a702a_2	52	KB	conda-forge
libgfortran-ng-14.2.0	h69a702a_2	53	ΚB	conda-forge
libgfortran5-14.2.0	hf1ad2bd_2	1.4	MB	conda-forge
libgomp-14.2.0	h767d61c_2	449	ΚB	conda-forge
libklu-2.3.5	hf24d653_7100102	142	ΚB	conda-forge
liblapack-3.9.0	31_h7ac8fdf_openblas		16	KB conda-forge
libld1-3.3.2	h32481e8_7100102	24	ΚB	conda-forge
libopenblas-0.3.29	pthreads_h94d23a6_0	5	.6 1	MB conda-forge
libparu-1.0.0	h17147ab_7100102	91	ΚB	conda-forge
librbio-4.3.4	h32481e8_7100102	47	ΚB	conda-forge
libspex-3.2.3	had10066_7100102	79	ΚB	conda-forge
libspqr-4.3.4	h852d39f_7100102	213	ΚB	conda-forge
libstdcxx-14.2.0	h8f9b012_2	3.7	MB	conda-forge
libstdcxx-ng-14.2.0	h4852527_2	53	ΚB	conda-forge
libsuitesparseconfig-7.10.1	h92d6892_7100102	42	ΚB	conda-forge
libumfpack-6.3.5	heb53515_7100102	424	ΚB	conda-forge
metis-5.1.0	hd0bcaf9_1007	3.7	MB	conda-forge
mpfr-4.2.1	h90cbb55_3	620	KB	conda-forge

```
numpy-2.2.4
                           py311h5d046bc_0
                                                 8.6 MB conda-forge
openssl-3.4.1
                              h7b32b05_0
                                                2.8 MB conda-forge
pyfmi-2.16.3
                           py311h9f3472d 0
                                                5.2 MB conda-forge
python_abi-3.11
                                   2_cp311
                                                  5 KB conda-forge
scipy-1.15.2
                          py311h8f841c2_0
                                                 16.4 MB conda-forge
suitesparse-7.10.1
                          ha0f6916_7100102
                                                 12 KB conda-forge
                                                907 KB conda-forge
sundials-7.1.1
                           ha52427a 0
                                    Total:
                                                56.1 MB
```

The following NEW packages will be INSTALLED:

```
_x86_64-microarch~ conda-forge/noarch::_x86_64-microarch-level-3-2_broadwell
                     conda-forge/linux-64::assimulo-3.6.0-py311h083bc19_0
  assimulo
 fmilib
                     conda-forge/linux-64::fmilib-2.4.1-hac33072 1
                     conda-forge/linux-64::gmp-6.3.0-hac33072_2
  gmp
                     conda-forge/linux-64::libamd-3.3.3-haaf9dc3_7100102
 libamd
                     conda-forge/linux-64::libblas-3.9.0-31_h59b9bed_openblas
 libblas
 libbtf
                     conda-forge/linux-64::libbtf-2.3.2-h32481e8_7100102
 libcamd
                     conda-forge/linux-64::libcamd-3.3.3-h32481e8_7100102
 libcblas
                     conda-forge/linux-64::libcblas-3.9.0-31_he106b2a_openblas
                     conda-forge/linux-64::libccolamd-3.3.4-h32481e8_7100102
 libccolamd
 libcholmod
                     conda-forge/linux-64::libcholmod-5.3.1-h59ddab4_7100102
                     conda-forge/linux-64::libcolamd-3.3.4-h32481e8_7100102
 libcolamd
 libcxsparse
                     conda-forge/linux-64::libcxsparse-4.4.1-h32481e8_7100102
 libgcc
                     conda-forge/linux-64::libgcc-14.2.0-h767d61c_2
                     conda-forge/linux-64::libgfortran-14.2.0-h69a702a_2
 libgfortran
 libgfortran-ng
                     conda-forge/linux-64::libgfortran-ng-14.2.0-h69a702a_2
 libgfortran5
                     conda-forge/linux-64::libgfortran5-14.2.0-hf1ad2bd_2
 libklu
                     conda-forge/linux-64::libklu-2.3.5-hf24d653_7100102
 liblapack
                     conda-forge/linux-64::liblapack-3.9.0-31 h7ac8fdf openblas
 libldl
                     conda-forge/linux-64::libldl-3.3.2-h32481e8 7100102
                     conda-forge/linux-64::libopenblas-0.3.29-pthreads_h94d23a6_0
 libopenblas
                     conda-forge/linux-64::libparu-1.0.0-h17147ab_7100102
 libparu
 librbio
                     conda-forge/linux-64::librbio-4.3.4-h32481e8_7100102
                     conda-forge/linux-64::libspex-3.2.3-had10066_7100102
 libspex
 libspqr
                     conda-forge/linux-64::libspqr-4.3.4-h852d39f 7100102
                     conda-forge/linux-64::libstdcxx-14.2.0-h8f9b012 2
  libstdcxx
 libsuitesparsecon~ conda-forge/linux-64::libsuitesparseconfig-7.10.1-h92d6892_7100
102
 libumfpack
                     conda-forge/linux-64::libumfpack-6.3.5-heb53515_7100102
 metis
                     conda-forge/linux-64::metis-5.1.0-hd0bcaf9_1007
  mpfr
                     conda-forge/linux-64::mpfr-4.2.1-h90cbb55_3
                     conda-forge/linux-64::numpy-2.2.4-py311h5d046bc 0
  numpy
  pyfmi
                     conda-forge/linux-64::pyfmi-2.16.3-py311h9f3472d_0
                     conda-forge/linux-64::python_abi-3.11-2_cp311
  python_abi
  scipy
                     conda-forge/linux-64::scipy-1.15.2-py311h8f841c2_0
  suitesparse
                     conda-forge/linux-64::suitesparse-7.10.1-ha0f6916_7100102
  sundials
                     conda-forge/linux-64::sundials-7.1.1-ha52427a_0
```

The following packages will be UPDATED:

```
libgomp
                     pkgs/main::libgomp-11.2.0-h1234567_1 --> conda-forge::libgomp
-14.2.0-h767d61c_2
                   pkgs/main::libstdcxx-ng-11.2.0-h12345~ --> conda-forge::libstdc
 libstdcxx-ng
xx-ng-14.2.0-h4852527_2
 openssl
                     pkgs/main::openssl-3.0.16-h5eee18b_0 --> conda-forge::openssl
-3.4.1-h7b32b05_0
The following packages will be SUPERSEDED by a higher-priority channel:
 certifi
                   pkgs/main/linux-64::certifi-2025.1.31~ --> conda-forge/noarch::
certifi-2025.1.31-pyhd8ed1ab_0
Downloading and Extracting Packages:
                   16.4 MB
scipy-1.15.2
                               | :
                                   0% 0/1 [00:00<?, ?it/s]
numpy-2.2.4
                   8.6 MB
                               | : 0% 0/1 [00:00<?, ?it/s]
libopenblas-0.3.29 | 5.6 MB
                               | : 0% 0/1 [00:00<?, ?it/s]
pyfmi-2.16.3
                   5.2 MB
                               |:
                                    0% 0/1 [00:00<?, ?it/s]
metis-5.1.0
                   3.7 MB
                               : 0% 0/1 [00:00<?, ?it/s]
libstdcxx-14.2.0
                   | 3.7 MB | : 0% 0/1 [00:00<?, ?it/s]
openssl-3.4.1
             | 2.8 MB | : 0% 0/1 [00:00<?, ?it/s]
libgfortran5-14.2.0 | 1.4 MB | : 0% 0/1 [00:00<?, ?it/s]
```

assimulo-3.6.0 | 1.1 MB | : 0% 0/1 [00:00<?, ?it/s]

libcholmod-5.3.1 | 1.1 MB | : 0% 0/1 [00:00<?, ?it/s]

sundials-7.1.1 | 907 KB | : 0% 0/1 [00:00<?, ?it/s]

libgcc-14.2.0 | 828 KB | : 0% 0/1 [00:00<?, ?it/s]

mpfr-4.2.1 | 620 KB | : 0% 0/1 [00:00<?, ?it/s]

gmp-6.3.0 | 449 KB | : 0% 0/1 [00:00<?, ?it/s]

libgomp-14.2.0 | 449 KB | : 0% 0/1 [00:00<?, ?it/s]

libumfpack-6.3.5 | 424 KB | : 0% 0/1 [00:00<?, ?it/s]

libspqr-4.3.4 | 213 KB | : 0% 0/1 [00:00<?, ?it/s]

... (more hidden) ...

libopenblas-0.3.29 | 5.6 MB | : 82% 0.8192985372565079/1 [00:00<00:00, 4.44i

pyfmi-2.16.3 | 5.2 MB | : 90% 0.8951859169945998/1 [00:00<00:00, 4.92i t/s]

metis-5.1.0 | 3.7 MB | : 100% 1.0/1 [00:00<00:00, 4.07it/s]

metis-5.1.0 | 3.7 MB | : 100% 1.0/1 [00:00<00:00, 4.07it/s]

scipy-1.15.2 | 16.4 MB | : 58% 0.5765280845379718/1 [00:00<00:00, 2.13i t/s]

pyfmi-2.16.3 | 5.2 MB | : 100% 1.0/1 [00:00<00:00, 4.92it/s]

libopenblas-0.3.29 | 5.6 MB | : 100% 1.0/1 [00:00<00:00, 4.44it/s]

openssl-3.4.1 | 2.8 MB | : 1% 0.0055741049077571376/1 [00:00<01:03, 64.2 8s/it]

libgfortran5-14.2.0 | 1.4 MB | : 1% 0.011206734985068174/1 [00:00<00:31, 32.27 s/it]

scipy-1.15.2 | 16.4 MB | : 79% 0.78903347768172/1 [00:00<00:00, 1.98it/s]

libgfortran5-14.2.0 | 1.4 MB | : 100% 1.0/1 [00:00<00:00, 32.27s/it] numpy-2.2.4 | 8.6 MB | : 100% 1.0/1 [00:00<00:00, 4.38it/s]

conda-25.1.1 | 1.1 MB | : 1% 0.013622478419712683/1 [00:00<00:34, 34.94 s/it]

libstdcxx-14.2.0 | 3.7 MB | : 100% 1.0/1 [00:00<00:00, 3.33it/s]

assimulo-3.6.0 | 1.1 MB | : 1% 0.014703493605362324/1 [00:00<00:33, 33.82 s/it]

openssl-3.4.1 | 2.8 MB | : 100% 1.0/1 [00:00<00:00, 2.46it/s]

openssl-3.4.1 | 2.8 MB | : 100% 1.0/1 [00:00<00:00, 2.46it/s]

scipy-1.15.2 | 16.4 MB | : 99% 0.9891506640502722/1 [00:00<00:00, 1.79i t/s]

libcholmod-5.3.1 | 1.1 MB | : 1% 0.014870549794649543/1 [00:00<00:35, 36.27 s/it]

sundials-7.1.1 | 907 KB | : 2% 0.01763373830085844/1 [00:00<00:30, 30.84
s/it]</pre>

assimulo-3.6.0 | 1.1 MB | : 100% 1.0/1 [00:00<00:00, 33.82s/it]

libgcc-14.2.0 | 828 KB | : 2% 0.01932337522187561/1 [00:00<00:28, 28.67 s/it]

sundials-7.1.1 | 907 KB | : 100% 1.0/1 [00:00<00:00, 30.84s/it]

libcholmod-5.3.1 | 1.1 MB | : 100% 1.0/1 [00:00<00:00, 36.27s/it]

libgcc-14.2.0 | 828 KB | : 100% 1.0/1 [00:00<00:00, 28.67s/it]

mpfr-4.2.1 | 620 KB | : 100% 1.0/1 [00:00<00:00, 22.82s/it]

libgomp-14.2.0 | 449 KB | : 4% 0.03562807972826631/1 [00:00<00:16, 17.31 s/it]

gmp-6.3.0 | 449 KB | : 4% 0.03561313321233331/1 [00:00<00:16, 17.40 s/it]

gmp-6.3.0 | 449 KB | : 100% 1.0/1 [00:00<00:00, 17.40s/it]

fmilib-2.4.1 | 383 KB | : 4% 0.04180391656566945/1 [00:00<00:14, 15.16 s/it]

libgomp-14.2.0 | 449 KB | : 100% 1.0/1 [00:00<00:00, 17.31s/it]

libumfpack-6.3.5 | 424 KB | : 4% 0.037731330084655984/1 [00:00<00:16, 17.18 s/it]

libumfpack-6.3.5 | 424 KB | : 100% 1.0/1 [00:00<00:00, 17.18s/it]

libspqr-4.3.4 | 213 KB | : 8% 0.07503068271326775/1 [00:00<00:08, 9.03 s/it]

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... (more hidden) ...

libspqr-4.3.4 | 213 KB | : 100% 1.0/1 [00:00<00:00, 9.03s/it]

metis-5.1.0 | 3.7 MB | : 100% 1.0/1 [00:00<00:00, 4.07it/s]

scipy-1.15.2 | 16.4 MB | : 100% 1.0/1 [00:00<00:00, 1.79it/s]

libgfortran5-14.2.0 | 1.4 MB | : 100% 1.0/1 [00:01<00:00, 1.11it/s]

libgfortran5-14.2.0 | 1.4 MB | : 100% 1.0/1 [00:01<00:00, 1.11it/s]

libopenblas-0.3.29 | 5.6 MB | : 100% 1.0/1 [00:01<00:00, 4.44it/s]

libstdcxx-14.2.0 | 3.7 MB | : 100% 1.0/1 [00:01<00:00, 3.33it/s]

openssl-3.4.1 | 2.8 MB | : 100% 1.0/1 [00:01<00:00, 2.46it/s]

conda-25.1.1 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.00s/it]

conda-25.1.1 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.00s/it]

assimulo-3.6.0 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.15s/it]

assimulo-3.6.0 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.15s/it] numpy-2.2.4 | 8.6 MB | : 100% 1.0/1 [00:02<00:00, 4.38it/s]

libcholmod-5.3.1 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.25s/it]

libcholmod-5.3.1 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.25s/it]

sundials-7.1.1 | 907 KB | : 100% 1.0/1 [00:02<00:00, 2.33s/it]

sundials-7.1.1 | 907 KB | : 100% 1.0/1 [00:02<00:00, 2.33s/it]

libgcc-14.2.0 | 828 KB | : 100% 1.0/1 [00:02<00:00, 2.35s/it]

libgcc-14.2.0 | 828 KB | : 100% 1.0/1 [00:02<00:00, 2.35s/it]

gmp-6.3.0 | 449 KB | : 100% 1.0/1 [00:02<00:00, 2.40s/it]

mpfr-4.2.1 | 620 KB | : 100% 1.0/1 [00:02<00:00, 2.42s/it]

mpfr-4.2.1 | 620 KB | : 100% 1.0/1 [00:02<00:00, 2.42s/it]

libumfpack-6.3.5 | 424 KB | : 100% 1.0/1 [00:02<00:00, 2.44s/it]

libumfpack-6.3.5 | 424 KB | : 100% 1.0/1 [00:02<00:00, 2.44s/it]

libgomp-14.2.0 | 449 KB | : 100% 1.0/1 [00:02<00:00, 2.48s/it]

libgomp-14.2.0 | 449 KB | : 100% 1.0/1 [00:02<00:00, 2.48s/it]

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... (more hidden) ...

fmilib-2.4.1 | 383 KB | : 100% 1.0/1 [00:02<00:00, 2.53s/it]

fmilib-2.4.1 | 383 KB | : 100% 1.0/1 [00:02<00:00, 2.53s/it]

libspqr-4.3.4 | 213 KB | : 100% 1.0/1 [00:02<00:00, 2.56s/it]

scipy-1.15.2 | 16.4 MB | : 100% 1.0/1 [00:03<00:00, 1.79it/s]

Preparing transaction: - 22\ 22done

Verifying transaction: / 22- 22\ 22| 22done

Executing transaction: - 22\ 22| 22/ 22- 22\ 22| 22/ 22- 22\ 22| 22/ 22- 22\ 22| 22/ 22- 22\ 22| 22/ 22done

Now specific installation and the run simulations. Start with connecting to Github. Then upload the four files:

- FMU BPL_TEST2_Batch_linux_om_me.fmu
- Setup-file BPL_TEST2_Batch_explore.py

Cloning into 'BPL_TEST2_Batch_calibration'...

In [10]: %cd BPL_TEST2_Batch_calibration

/content/BPL_TEST2_Batch_calibration

BPL_TEST2_Batch_calibration - demo

Author: Jan Peter Axelsson

This notebook shows the possibilities for calibration of the model BPL_TEST2_Batch using scipy.optimize.minimize() routine. There are several different methods to choose between. In this notebook we work with simulated data.

The text-book model of batch cultivation we simulate is the following where S is substrate, X is cell concentration, and V is volume of the broth

$$\frac{d(VS)}{dt} = -q_S(S) \cdot VX$$

$$\frac{d(VX)}{dt} = \mu(S) \cdot VX$$

and where specific cell growth rate μ and substrate uptake rate q_S are

$$\mu(S) = Y \cdot q_S(S)$$

$$q_S(S) = q_S^{max} rac{S}{K_s + S}$$

where Y is the yield, q_S^{max} is the maximal specific substrate uptake rate and K_s is the corresponding saturation constant.

The parameter estimation is done with optimization methods that only require evaluation of the missmatch between simulation with given parameters and data. At start the allowed range for each parameter is given. The method used for optimization is Nelder-Mead but can easily be changed [1].

In the near future the FMU may provide first derivative gradient information, that will make it possible to choose corresponding method of minimize() for improved performance. This possibility is related to the upgrade to the FMI-standard ver 3.0 for the Modelica compiler.

The Python package PyFMI [2] that is the base for FMU-explore has a simplified built-in functionality for parameter estimation that also use scipy.optimize.minimize(). However, there is estimatation functionally but the purpose seems to only address smaller examples. There is for instance no support to handle models that takes sub-models from libraries and necessary changes of default parameters not to be estimated. Therefore we here define a Python function evaluate() that facilitate the formulation of the parameter estimation and also bring flexibility to choice of optimization method, default Nelder-Mead.

```
In [11]: run -i BPL_TEST2_Batch_explore.py
Linux - run FMU pre-compiled OpenModelica
```

·

Model for the process has been setup. Key commands:

- par() change of parameters and initial values
- init()change initial values only
- simu() simulate and plot
- newplot() make a new plot
- show() show plot from previous simulation
- disp()
 display parameters and initial values from the last simulation
- describe() describe culture, broth, parameters, variables with values/units

Note that both disp() and describe() takes values from the last simulation and the command process_diagram() brings up the main configuration

Brief information about a command by help(), eg help(simu)
Key system information is listed with the command system_info()

```
In [12]: # Adjust the size of diagrams
plt.rcParams['figure.figsize'] = [15/2.54, 12/2.54]
```

1 Generate data later used for parameter estimation

```
In [13]: import pandas as pd
```

```
In [14]: # Data generated
    simulationTime = 6.0
    par(Y=0.50, qSmax=1.00, Ks=0.1)
    init(V_start=1.0, VS_start=10, VX_start=1.0)
    newplot(plotType='Demo_2')
    simu(simulationTime, options=opts_data)
```

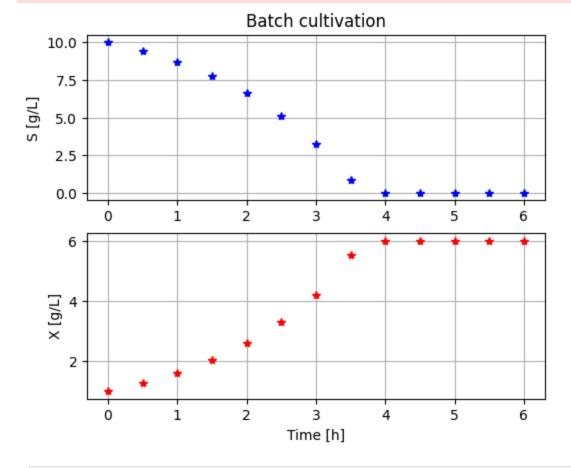
```
Could not find cannot import name 'dopri5' from 'assimulo.lib' (/usr/local/lib/pytho n3.11/site-packages/assimulo/lib/__init__.py)

Could not find cannot import name 'rodas' from 'assimulo.lib' (/usr/local/lib/python 3.11/site-packages/assimulo/lib/__init__.py)

Could not find cannot import name 'odassl' from 'assimulo.lib' (/usr/local/lib/pytho n3.11/site-packages/assimulo/lib/__init__.py)

Could not find ODEPACK functions.

Could not find GLIMDA.
```



```
In [15]: # Store data in a DataFrame for Later use
data = pd.DataFrame(data={'time':sim_res['time'], 'X':sim_res['bioreactor.c[1]'], '
data
```

Out[15]:		time	Х	S
	0	0.0	1.000000	1.000000e+01
	1	0.5	1.269848	9.438455e+00
	2	1.0	1.615795	8.719839e+00
	3	1.5	2.050445	7.800734e+00
	4	2.0	2.601038	6.626389e+00
	5	2.5	3.297304	5.128962e+00
	6	3.0	4.195962	3.229259e+00
	7	3.5	5.524388	8.813998e-01
	8	4.0	6.000000	-2.037810e-08
	9	4.5	6.000000	2.960320e-10
	10	5.0	6.000000	1.200938e-10
	11	5.5	6.000000	2.363337e-10
	12	6.0	6.000000	-1.553435e-10

2 Simulation with initial guess of parameters compared with data

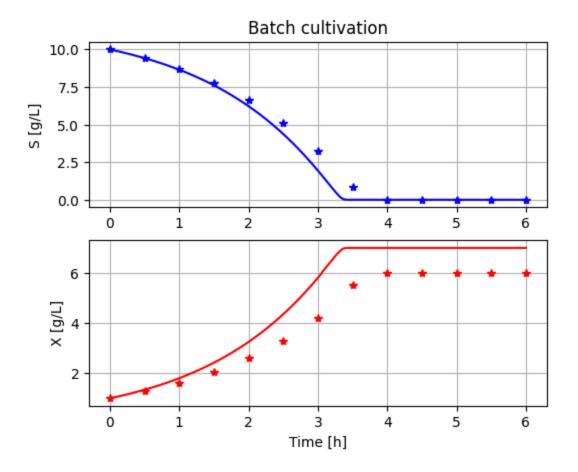
Here we define the parameters that should be estimated and specify allowed ranges. Nominal parameters are chosen as the mid-point of the allowed parameter range.

Simulation with these nominal parameter set and compare with data give an idea of who well the model fit data.

```
In [16]: # Parameters to be estimated using parDict names and their bounds
    parEstim = ['Y', 'qSmax', 'Ks']
    parBounds = [(0.4, 0.8), (0.7, 1.3), (0.05, 0.20)]
    parEstim_0 = [np.mean(parBounds[k]) for k in range(len(parBounds))]

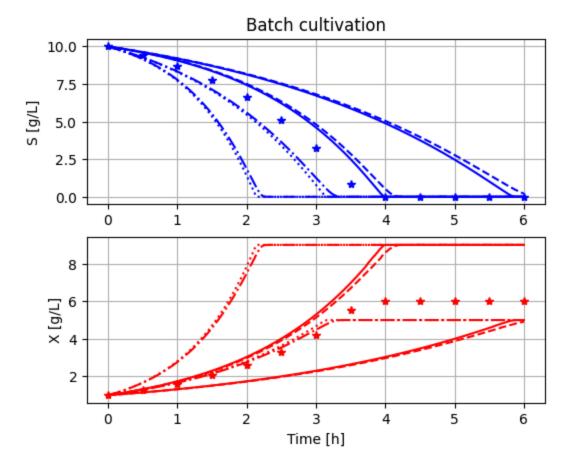
In [17]: # Simulation with nominal parameters
    newplot(plotType='Demo_1')
    par(Y=parEstim_0[0], qSmax=parEstim_0[1], Ks=parEstim_0[2])
    simu(simulationTime)

# Show data
    ax1.plot(data['time'], data['S'],'b*')
    ax2.plot(data['time'], data['X'],'r*')
    plt.show()
```



```
In [18]: # Simulation over the parameter ranges given
newplot(plotType='Demo_1')
for Y_value in parBounds [0]:
    for qSmax_value in parBounds[1]:
        for Ks_value in parBounds[2]:
            par(Y=Y_value, qSmax=qSmax_value, Ks=Ks_value)
            simu(simulationTime)

# Show data
ax1.plot(data['time'], data['S'],'b*')
ax2.plot(data['time'], data['X'],'r*')
plt.show()
```



Simulation over the different parameter combinations of the parameter bounds shows that data is "covered" and we have good hope to find a parameter combination that fits data well.

3 Parameter estimation

Here we use the scipy.optimize.minimize() procedure which contain a family of different methods [1]. The default method is Nelder-Mead and is robust for fitting a model to data. Further we have chosen to work with bounds for the parameters to be estimated and the initial guess is chosen as the middle point in parameter space.

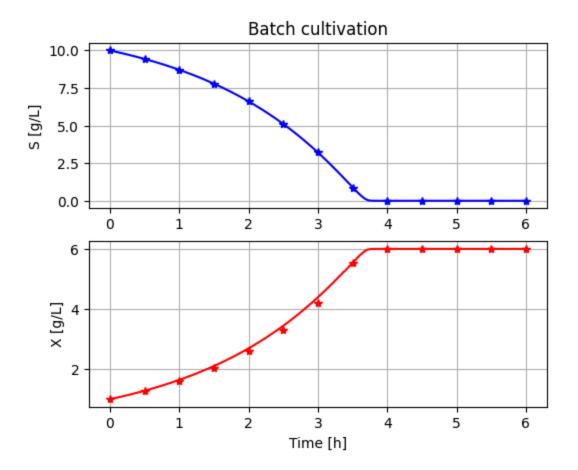
```
# Update parameters and simulate
             for i, p in enumerate(parEstim): par(**{p:x[i]})
             simu(simulationTime, options=opts_fast)
             # Calculate loss function V
             V={}
             V['X'] = np.linalg.norm(data['X'] - np.interp(data['time'], sim_res['time'], si
             V['S'] = np.linalg.norm(data['S'] - np.interp(data['time'], sim_res['time'], si
             return V['X'] + V['S']
In [22]: import time
In [23]: # Run minimize()
         start_time = time.time()
         result = scipy.optimize.minimize(objective, x0=parEstim_0, args=extra_args,
                                           method='Nelder-Mead', bounds=parBounds, options={"
         print('CPU-time =', time.time()-start_time)
        Optimization terminated successfully.
                 Current function value: 0.148311
                 Iterations: 66
                 Function evaluations: 122
        CPU-time = 0.5640532970428467
In [24]: result.x
Out[24]: array([0.49997276, 1.00731527, 0.14380564])
```

The estimated parameters result.x are very close to the original values and no surprise.

4 Simulation with estimated parameters compared with data

```
In [25]: newplot(plotType='Demo_1')
    par(Y=result.x[0], qSmax=result.x[1], Ks=result.x[2])
    simu(simulationTime)

# Show data
    ax1.plot(data['time'], data['S'],'b*')
    ax2.plot(data['time'], data['X'],'r*')
    plt.show()
```



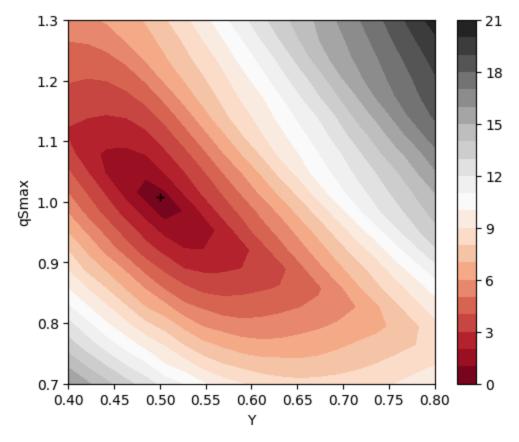
```
In [26]: # The estimated parameters are
for i in range(len(parEstim)): print(parEstim[i],':', result.x[i])
```

Y : 0.4999727558733863 qSmax : 1.0073152667279195 Ks : 0.14380564144282715

5 Analysis of the loss function

The problem is small and analysis of the loss function brings some insight. From the diagram above showing parameter sweep over combinations min- and max-parameters we see that the parameter K_s has little influence. Let use set that a fixed value and then plot the loss function in the parameters Y and qSmax. We do this by go through all the parametera combinations and evaluate each of them.

```
# Contour plot
plt.figure()
plt.clf
plt.subplot(1,1,1)
plt.contourf(Y, qSmax, V, 20, cmap='RdGy')
plt.plot(result.x[0], result.x[1],'k+')
plt.colorbar()
plt.ylabel('qSmax')
plt.xlabel('Y')
plt.show()
```



We see the following in the contour diagram of the loss function simplified:

- The minima is unique in the range of parmaters we study. This is good news.
- The contour plot is ellipsoid and rather narrow. The more narrow the ellipsoid the more difficult and more time it takes to converge to the minima.
- The direction of the ellipsoid axis indicate the correlation you may get between the two parameters during the minimization process.

Note that the form of the contour plot change with the parameters (and initial values) of the actual proces. You can see the impact by changing the parameters in "cell # 4" where data is generated and then just choose to run that cell and the cells below. No need to restart the notebook.

6 Summary

A choice was made to work with allowed ranges of parameters to be estimated and a start value was defined as the center point in this parameter space. There are only three methods available in optimize.minimize() that can handle bounds on parameters.

An evaluate() function was created that define how the difference beween simulation and data is measured. The function is rather transparent and easy to modify and you may want to change weight on the loss in S and X, for instance. Here they have so far equal weight.

The FMU-explore workspace dictionaries partDict[] and parLocation[] are useful also here and simplify the code for the evaluation() function. But we also use the detailed PyFMI-functions to administrate and set parameters of the actual simulation.

The call optimize.minimize() has several parameters and can easily be modified, for instance change of method. For fitting a model to data Nelder-Mead is ao a robust and good choice, but can be somewhat slow.

The estimated parameters were close to perfect!

The contour plot of the simplified loss function shows that the minima is unique and should not be difficult too difficut to obtain. More narrow elliptical contour plots would indicate difficulties. Multiple local minima would also be a problem.

7 References

- [1] Scipy Reference guide on optimize.minimize() here
- [2] Andersson, C., Åkesson, J., Fuhrer C.: "PyFMI: A Python package for simulation of coupled dynamic models with the functional mock-up interface", Centre for Mathematical Sciences, Lund University, Report LUTFNA-5008-2016, 2016.

Appendix

```
In [28]: describe('parts')
        ['bioreactor', 'bioreactor.culture']
In [29]: describe('MSL')
        MSL: 3.2.3 - used components: none
In [30]: system_info()
```

System information

-OS: Linux

-Python: 3.11.11 -Scipy: 1.14.1 -PyFMI: 2.16.3

-FMU by: OpenModelica Compiler OpenModelica 1.25.0~dev-133-ga5470be

-FMI: 2.0

-Type: FMUModelME2

-Name: BPL.Examples_TEST2.Batch -Generated: 2024-11-06T21:35:34Z

-MSL: 3.2.3

-Description: Bioprocess Library version 2.3.0

-Interaction: FMU-explore version 1.0.0

In [30]: