

▼ BPL_TEST2_Chemostat script with PyFMI ver 2.9.8

The key library PyFMI v2.9.8 is installed.

After the installation a small application BPL_TEST2_Chemostat is loaded and run. You can continue with this example if you like.

```
lslsb_release -a # Actual VM Ubuntu version used by Google

No LSB modules are available.
Distributor ID: Ubuntu
Description:   Ubuntu 20.04.5 LTS
Release:       20.04
Codename:      focal

%env PYTHONPATH=

env: PYTHONPATH=

!wget https://repo.anaconda.com/miniconda/Miniconda3-py38_22.11.1-1-Linux-x86_64.sh
!chmod +x Miniconda3-py38_22.11.1-1-Linux-x86_64.sh
!bash ./Miniconda3-py38_22.11.1-1-Linux-x86_64.sh -b -f -p /usr/local
import sys
sys.path.append('/usr/local/lib/python3.8/site-packages/')

--2023-01-21 09:37:30--  https://repo.anaconda.com/miniconda/Miniconda3-py38\_22.11.1-1-Linux-x86\_64.sh
Resolving repo.anaconda.com (repo.anaconda.com)... 104.16.130.3, 104.16.131.3, 2606:4700::6810:8303, ...
Connecting to repo.anaconda.com (repo.anaconda.com)|104.16.130.3|:443... connected.
HTTP request sent, awaiting response... 200 OK
Length: 64630241 (62M) [application/x-sh]
Saving to: 'Miniconda3-py38_22.11.1-1-Linux-x86_64.sh'

Miniconda3-py38_22. 100%[=====] 61.64M  195MB/s   in 0.3s

2023-01-21 09:37:30 (195 MB/s) - 'Miniconda3-py38_22.11.1-1-Linux-x86_64.sh' saved [64630241/64630241]

PREFIX=/usr/local
Unpacking payload ...

Installing base environment...

Downloading and Extracting Packages

Downloading and Extracting Packages

Preparing transaction: done
Executing transaction: done
installation finished.

!conda update -n base -c defaults conda --yes
```

```

numpy -> libgcc-ng[version='>=11.2.0'] -> _openmp_mutex[version='>=4.5']
python=3.8 -> libgcc-ng[version='>=11.2.0'] -> _openmp_mutex[version='>=4.5']
openssl -> libgcc-ng[version='>=7.5.0'] -> _openmp_mutex[version='>=4.5']
scipy -> libgcc-ng[version='>=11.2.0'] -> _openmp_mutex[version='>=4.5']
suitesparse -> libgcc-ng[version='>=11.2.0'] -> _openmp_mutex[version='>=4.5']

```

Package cryptography conflicts for:

```
conda-content-trust -> cryptography
```

```
conda -> pyopenssl[version='>=16.2.0'] -> cryptography[version='>=1.9|>=2.1.4|>=2.2.1|>=2.8|>=3.3|>=35.0']
```

```
conda[version='>=22.11.1'] -> pyopenssl[version='>=16.2.0'] -> cryptography[version='>=1.9|>=2.1.4|>=2.2.1|>=2.8|>=3.3|>=
```

Package libhwloc conflicts for:

```
tbb -> libhwloc[version='>=2.8.0,<2.8.1.0a0']
```

```
suitesparse -> tbb[version='>=2020.3'] -> libhwloc[version='>=2.8.0,<2.8.1.0a0']The following specifications were found t
```

```

- feature:/linux-64::__glibc==2.31=0
- feature:|@/linux-64::__glibc==2.31=0
- assimulo -> libgfortran-ng -> __glibc[version='>=2.17']
- libopenblas -> libgcc-ng[version='>=11.2.0'] -> __glibc[version='>=2.17']
- openssl -> libgcc-ng[version='>=7.5.0'] -> __glibc[version='>=2.17']
- python=3.8 -> libgcc-ng[version='>=11.2.0'] -> __glibc[version='>=2.17']
- scipy -> libgcc-ng[version='>=11.2.0'] -> __glibc[version='>=2.17']
- suitesparse -> libgcc-ng[version='>=11.2.0'] -> __glibc[version='>=2.17']
- tbb -> libgcc-ng[version='>=11.2.0'] -> __glibc[version='>=2.17']

```

Your installed version is: 2.31

```
!conda --version
!python --version
```

```
conda 22.11.1
Python 3.8.15
```

```
!conda install -c conda-forge pyfmi --yes # Install the key package
```

Collecting package metadata (current_repodata.json): done

Solving environment: |

The environment is inconsistent, please check the package plan carefully

The following packages are causing the inconsistency:

```

- conda-forge/linux-64::tbb==2021.7.0=h924138e_1
- conda-forge/linux-64::numpy==1.24.1=py38hab0fcb9_0
- conda-forge/linux-64::suitesparse==5.10.1=h9e50725_1
- conda-forge/linux-64::scipy==1.10.0=py38h10c12cc_0
- conda-forge/linux-64::libcblas==3.9.0=16_linux64_openblas
- conda-forge/linux-64::libopenblas==0.3.21=threads_h78a6416_3
- conda-forge/linux-64::pyfmi==2.9.8=py38h26c90d9_1
- conda-forge/linux-64::assimulo==3.3=py38h71f17ff_1
- conda-forge/linux-64::libblas==3.9.0=16_linux64_openblas
- conda-forge/linux-64::liblapack==3.9.0=16_linux64_openblas
- conda-forge/linux-64::libhwloc==2.8.0=h32351e8_1
- conda-forge/linux-64::fmlib==2.4.1=h27087fc_0
- conda-forge/linux-64::sundials==6.4.1=h89a52a3_done

```

Package Plan

environment location: /usr/local

added / updated specs:

```
- pyfmi
```

The following packages will be REMOVED:

```
libgomp-11.2.0-h1234567_1
```

The following packages will be UPDATED:

```

ca-certificates  pkgs/main::ca-certificates-2022.10.11~ --> conda-forge::ca-certificates-2022.12.7-ha878542_0
libgcc-ng        pkgs/main::libgcc-ng-11.2.0-h1234567_1 --> conda-forge::libgcc-ng-12.2.0-h65d4601_19
libstdcxx-ng     pkgs/main::libstdcxx-ng-11.2.0-h12345~ --> conda-forge::libstdcxx-ng-12.2.0-h46fd767_19
openssl          pkgs/main::openssl-1.1.1s-h7f8727e_0 --> conda-forge::openssl-1.1.1s-h0b41bf4_1

```

The following packages will be SUPERSEDED by a higher-priority channel:

```

_libgcc_mutex    pkgs/main::_libgcc_mutex-0.1-main --> conda-forge::_libgcc_mutex-0.1-conda_forge
_openmp_mutex    pkgs/main::_openmp_mutex-5.1-1_gnu --> conda-forge::_openmp_mutex-4.5-2_kmp_llvm
certifi          pkgs/main/linux-64::certifi-2022.12.7~ --> conda-forge/noarch::certifi-2022.12.7-pyhd8ed1ab_0
conda            pkgs/main::conda-22.11.1-py38h06a4308~ --> conda-forge::conda-22.11.1-py38h578d9bd_1

```

Downloading and Extracting Packages

Preparing transaction: done

Verifying transaction: done

Executing transaction: done

▼ Now specific installation run a simulation and notebook for that

Start with connecting to Github. Then upload the two files:

- FMU - BPL_TEST2_Chemostat_linux_om_me.fmu
- Setup-file - BPL_TEST2_Chemostat_explore.py

```
%%bash
git clone https://github.com/janpeter19/BPL_TEST2_Chemostat

Cloning into 'BPL_TEST2_Chemostat'...

%cd BPL_TEST2_Chemostat

/content/BPL_TEST2_Chemostat/BPL_TEST2_Chemostat
```

▼ BPL_TEST2_Chemostat - demo

```
run -i BPL_TEST2_Chemostat_explore_me.py
```

Linux - run FMU pre-compiled OpenModelica 1.21.0

Model for bioreactor has been setup. Key commands:

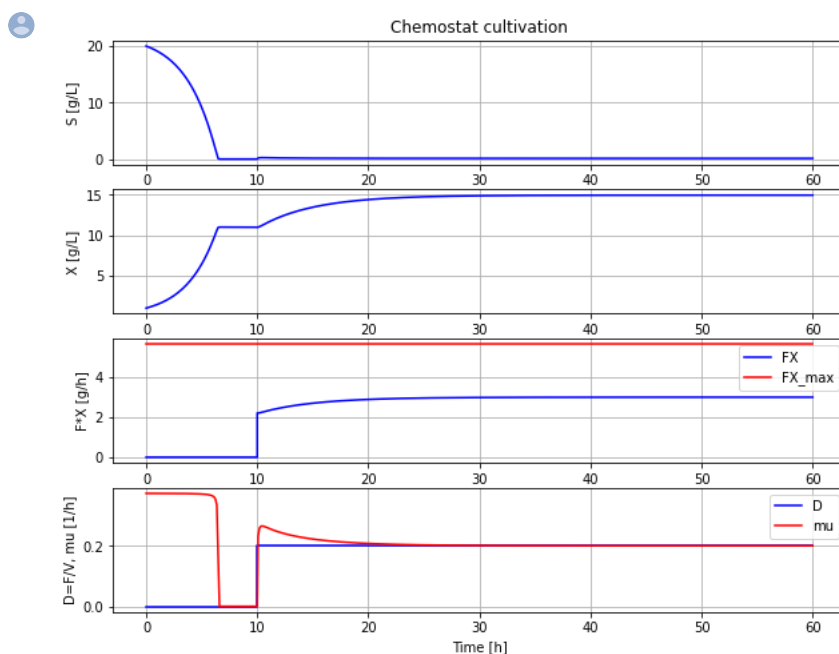
- par() - change of parameters and initial values
- init() - change initial values only
- simu() - simulate and plot
- newplot() - make a new plot
- show() - show plot from previous simulation
- disp() - display parameters and initial values from the last simulation
- describe() - describe culture, broth,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

Note that both disp() and describe() takes values from the last simulation

Brief information about a command by help(), eg help(simu)
Key system information is listed with the command system_info()
<Figure size 708.661x566.929 with 0 Axes>

```
%matplotlib inline
plt.rcParams['figure.figsize'] = [25/2.54, 20/2.54]
```

```
newplot()
par(Y=0.50, qSmax=0.75, Ks=0.1)      # Culture parameters
init(V_0=1.0, VX_0=1.0, VS_0=20)    # Bioreactor startup
par(S_in=30, t0=0, F0=0, t1=10, F1=0.2) # Substrate feeding
simu(60)
```



The maximal biomass productivity FX_{max} [g/h] marked red in the diagram above
can be calculated for CSTR from the FMU and is
cstrProdMax(model)

5.625

```
describe('cstrProdMax')
```

Calculate from the model maximal chemostat productivity FX_max : 5.625 [g/h]

```
describe('parts')
```

['bioreactor', 'bioreactor.culture', 'D', 'dosagescheme', 'feedtank', 'harvesttank']

```
describe('MSL')
```

MSL: 3.2.3 - used components: RealInput, RealOutput, CombiTimeTable, Types

```
system_info()
```

System information

- OS: Linux
- Python: 3.8.10
- Scipy: not installed in the notebook
- PyFMI: 2.9.8
- FMU by: OpenModelica Compiler OpenModelica 1.21.0~dev-185-g9d983b8
- FMI: 2.0
- Type: FMUModelME2
- Name: BPL_TEST2.Chemostat
- Generated: 2023-01-16T10:00:44Z
- MSL: 3.2.3
- Description: Bioprocess Library version 2.1.1-beta
- Interaction: FMU-explore version 0.9.6e

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✓ 0s completed at 10:41

