BPL_IEC_validation script with PyFMI

The key library PyFMI is installed.

After the installation a small application BPL_IEC_validation 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=
In [3]: !python --version
       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/')
       --2025-03-27 07:22:33-- https://repo.anaconda.com/miniconda/Miniconda3-py311_24.11.
       1-0-Linux-x86 64.sh
       Resolving repo.anaconda.com (repo.anaconda.com)... 104.16.191.158, 104.16.32.241, 26
       06:4700::6810:bf9e, ...
       Connecting to repo.anaconda.com (repo.anaconda.com) 104.16.191.158:443... connecte
       d.
       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
                                                                124MB/s
                                                                           in 1.1s
       2025-03-27 07:22:35 (124 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 \ 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	h06a4308 0	129 KB
certifi-2025.1.31	py311h06a4308 0	163 KB
openssl-3.0.16	h5eee18b_0	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, 11.14it/s]

certifi-2025.1.31 | 163 KB | : 69% 0.6888085428403262/1 [00:00<00:00, 6.71i

t/s]

ca-certificates-2025 | 129 KB | : 100% 1.0/1 [00:00<00:00, 9.83it/s]

openssl-3.0.16 | 5.2 MB | : 2% 0.017895563700972384/1 [00:00<00:05, 6.01

s/it]

certifi-2025.1.31 | 163 KB | : 100% 1.0/1 [00:00<00:00, 6.71it/s]

Preparing transaction: - 22done

Verifying transaction: | 22/22-22done

Executing transaction: | 22done

Channels:

- conda-forge
- defaults

Platform: linux-64

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

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-25.3.0	py311h38be061_0	1.1	MB	conda-forge
fmilib-2.4.1	hac33072_1	383	KB	conda-forge
gmp-6.3.0	hac33072_2	449	ΚB	conda-forge
libamd-3.3.3	haaf9dc3_7100102	49	ΚB	conda-forge
libblas-3.9.0	31_h59b9bed_openblas		16	KB conda-forge
libbtf-2.3.2	h32481e8_7100102	27	ΚB	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	ΚB	conda-forge
libgfortran-14.2.0	h69a702a_2	52	ΚB	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
libldl-3.3.2	h32481e8_7100102	24	KB	conda-forge
libopenblas-0.3.29	pthreads_h94d23a6_0	5	.6 1	MB conda-forge
libparu-1.0.0	h17147ab_7100102	91	KB	conda-forge
librbio-4.3.4	h32481e8_7100102	47	KB	conda-forge
libspex-3.2.3	had10066_7100102	79	KB	conda-forge
libspqr-4.3.4	h852d39f_7100102	213	KB	conda-forge
libstdcxx-14.2.0	h8f9b012_2	3.7	MB	conda-forge
libstdcxx-ng-14.2.0	h4852527_2	53	KB	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
                                                  5.2 MB conda-forge
pyfmi-2.16.3
                            py311h9f3472d_0
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
sundials-7.1.1
                                 ha52427a_0
                                                   907 KB
                                                           conda-forge
                                     Total:
                                                  56.1 MB
```

The following NEW packages will be INSTALLED:

```
_x86_64-microarch~ conda-forge/noarch::_x86_64-microarch-level-3-2_broadwell
  assimulo
                     conda-forge/linux-64::assimulo-3.6.0-py311h083bc19 0
                     conda-forge/linux-64::fmilib-2.4.1-hac33072_1
 fmilib
                     conda-forge/linux-64::gmp-6.3.0-hac33072_2
 gmp
                     conda-forge/linux-64::libamd-3.3.3-haaf9dc3_7100102
 libamd
 libblas
                     conda-forge/linux-64::libblas-3.9.0-31_h59b9bed_openblas
 libbtf
                     conda-forge/linux-64::libbtf-2.3.2-h32481e8_7100102
                     conda-forge/linux-64::libcamd-3.3.3-h32481e8_7100102
 libcamd
                     conda-forge/linux-64::libcblas-3.9.0-31_he106b2a_openblas
 libcblas
 libccolamd
                     conda-forge/linux-64::libccolamd-3.3.4-h32481e8_7100102
 libcholmod
                     conda-forge/linux-64::libcholmod-5.3.1-h59ddab4_7100102
 libcolamd
                     conda-forge/linux-64::libcolamd-3.3.4-h32481e8_7100102
 libcxsparse
                     conda-forge/linux-64::libcxsparse-4.4.1-h32481e8_7100102
                     conda-forge/linux-64::libgcc-14.2.0-h767d61c_2
 libgcc
 libgfortran
                     conda-forge/linux-64::libgfortran-14.2.0-h69a702a 2
 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
                     conda-forge/linux-64::libldl-3.3.2-h32481e8_7100102
 libldl
                     conda-forge/linux-64::libopenblas-0.3.29-pthreads h94d23a6 0
 libopenblas
 libparu
                     conda-forge/linux-64::libparu-1.0.0-h17147ab_7100102
 librbio
                     conda-forge/linux-64::librbio-4.3.4-h32481e8_7100102
 libspex
                     conda-forge/linux-64::libspex-3.2.3-had10066 7100102
                     conda-forge/linux-64::libspqr-4.3.4-h852d39f_7100102
 libspgr
                     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
                     conda-forge/linux-64::metis-5.1.0-hd0bcaf9_1007
 metis
  mpfr
                     conda-forge/linux-64::mpfr-4.2.1-h90cbb55 3
  numpy
                     conda-forge/linux-64::numpy-2.2.4-py311h5d046bc_0
                     conda-forge/linux-64::pyfmi-2.16.3-py311h9f3472d_0
  pyfmi
                     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
                     conda-forge/linux-64::sundials-7.1.1-ha52427a_0
  sundials
```

The following packages will be UPDATED:

```
libgomp
                     pkgs/main::libgomp-11.2.0-h1234567_1 --> conda-forge::libgomp
-14.2.0-h767d61c 2
 libstdcxx-ng
                    pkgs/main::libstdcxx-ng-11.2.0-h12345~ --> conda-forge::libstdc
xx-ng-14.2.0-h4852527_2
                     pkgs/main::openssl-3.0.16-h5eee18b_0 --> conda-forge::openssl
 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:
scipy-1.15.2
                    | 16.4 MB
                               | :
                                     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
                               1:
                                     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]
```

| : 0% 0/1 [00:00<?, ?it/s]

ng-14.2.0-h69a702a_2

conda-25.3.0

1.1 MB

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 | : 52% 0.5175973867127263/1 [00:00<00:00, 2.83i t/s] metis-5.1.0 3.7 MB | : 77% 0.7725229128648472/1 [00:00<00:00, 4.31i t/s] pyfmi-2.16.3 5.2 MB : 51% 0.5072720196302732/1 [00:00<00:00, 2.67i t/s] | : 100% 1.0/1 [00:00<00:00, 4.31it/s] metis-5.1.0 3.7 MB scipy-1.15.2 | 16.4 MB | : 32% 0.3192345592069761/1 [00:00<00:00, 1.26i t/s] scipy-1.15.2 | 16.4 MB | : 54% 0.5412693421777983/1 [00:00<00:00, 1.61i t/s] pyfmi-2.16.3 5.2 MB : 100% 1.0/1 [00:00<00:00, 2.48it/s] | 5.2 MB | : 100% 1.0/1 [00:00<00:00, 2.48it/s] pyfmi-2.16.3 libopenblas-0.3.29 5.6 MB | : 100% 1.0/1 [00:00<00:00, 2.34it/s] libopenblas-0.3.29 5.6 MB | : 100% 1.0/1 [00:00<00:00, 2.34it/s] openssl-3.4.1 2.8 MB | : 1% 0.0055741049077571376/1 [00:00<01:24, 85.3 4s/it] scipy-1.15.2 | 16.4 MB | : 70% 0.7042219082207621/1 [00:00<00:00, 1.48i t/s]

| 3.7 MB | : 100% 1.0/1 [00:00<00:00, 2.24it/s]

libstdcxx-14.2.0

libstdcxx-14.2.0 | 3.7 MB | : 100% 1.0/1 [00:00<00:00, 2.24it/s]

conda-25.3.0 | 1.1 MB | : 1% 0.01395284272989957/1 [00:00<00:42, 42.62 s/it]

libgfortran5-14.2.0 | 1.4 MB | : 100% 1.0/1 [00:00<00:00, 2.12it/s]

scipy-1.15.2 | 16.4 MB | : 86% 0.8557392064712374/1 [00:00<00:00, 1.48i

t/s]

numpy-2.2.4 | 8.6 MB | : 100% 1.0/1 [00:00<00:00, 3.25it/s]

assimulo-3.6.0 | 1.1 MB | : 1% 0.014703493605362324/1 [00:00<00:44, 44.89 s/it]

conda-25.3.0 | 1.1 MB | : 100% 1.0/1 [00:00<00:00, 42.62s/it]

openssl-3.4.1 | 2.8 MB | : 100% 1.0/1 [00:00<00:00, 1.90it/s]

openssl-3.4.1 | 2.8 MB | : 100% 1.0/1 [00:00<00:00, 1.90it/s]

libcholmod-5.3.1 | 1.1 MB | : 1% 0.014870549794649543/1 [00:00<00:46, 47.10 s/it]

assimulo-3.6.0 | 1.1 MB | : 100% 1.0/1 [00:00<00:00, 44.89s/it]

sundials-7.1.1 \mid 907 KB \mid : 2% 0.01763373830085844/1 [00:00<00:39, 40.39 s/it]

libgcc-14.2.0 | 828 KB | : 2% 0.01932337522187561/1 [00:00<00:37, 38.12 s/it]

sundials-7.1.1 | 907 KB | : 100% 1.0/1 [00:00<00:00, 40.39s/it]

mpfr-4.2.1 | 620 KB | : 3% 0.025811696239942908/1 [00:00<00:28, 29.07 s/it]

libgcc-14.2.0 | 828 KB | : 100% 1.0/1 [00:00<00:00, 38.12s/it]

mpfr-4.2.1 | 620 KB | : 100% 1.0/1 [00:00<00:00, 29.07s/it]

gmp-6.3.0 | 449 KB | : 4% 0.03561313321233331/1 [00:00<00:21, 22.25 s/it]

libgomp-14.2.0 | 449 KB | : 4% 0.03562807972826631/1 [00:00<00:21, 22.37 s/it]

libgomp-14.2.0 | 449 KB | : 100% 1.0/1 [00:00<00:00, 22.37s/it]

libumfpack-6.3.5 | 424 KB | : 4% 0.037731330084655984/1 [00:00<00:21, 22.05 s/it]

fmilib-2.4.1 | 383 KB | : 4% 0.04180391656566945/1 [00:00<00:19, 20.13 s/it]

... (more hidden) ...

... (more hidden) ...

libumfpack-6.3.5 | 424 KB | : 100% 1.0/1 [00:00<00:00, 22.05s/it]

metis-5.1.0 | 3.7 MB | : 100% 1.0/1 [00:00<00:00, 4.31it/s]

libspqr-4.3.4 | 213 KB | : 8% 0.07503068271326775/1 [00:00<00:10, 11.87 s/it]

libspqr-4.3.4 | 213 KB | : 100% 1.0/1 [00:00<00:00, 11.87s/it]

scipy-1.15.2 | 16.4 MB | : 100% 1.0/1 [00:01<00:00, 1.48it/s]

libstdcxx-14.2.0 | 3.7 MB | : 100% 1.0/1 [00:01<00:00, 2.24it/s]

libopenblas-0.3.29 | 5.6 MB | : 100% 1.0/1 [00:01<00:00, 2.34it/s]

libgfortran5-14.2.0 | 1.4 MB | : 100% 1.0/1 [00:01<00:00, 2.12it/s]

conda-25.3.0 | 1.1 MB | : 100% 1.0/1 [00:01<00:00, 1.78s/it]

conda-25.3.0 | 1.1 MB | : 100% 1.0/1 [00:01<00:00, 1.78s/it]

openssl-3.4.1 | 2.8 MB | : 100% 1.0/1 [00:02<00:00, 1.90it/s]

assimulo-3.6.0 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.23s/it]

assimulo-3.6.0 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.23s/it]

libcholmod-5.3.1 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.29s/it]

libcholmod-5.3.1 | 1.1 MB | : 100% 1.0/1 [00:02<00:00, 2.29s/it]

sundials-7.1.1 | 907 KB | : 100% 1.0/1 [00:02<00:00, 2.51s/it]

sundials-7.1.1 | 907 KB | : 100% 1.0/1 [00:02<00:00, 2.51s/it]

libgcc-14.2.0 | 828 KB | : 100% 1.0/1 [00:02<00:00, 2.63s/it]

libgcc-14.2.0 | 828 KB | : 100% 1.0/1 [00:02<00:00, 2.63s/it]

mpfr-4.2.1 | 620 KB | : 100% 1.0/1 [00:02<00:00, 2.71s/it]

mpfr-4.2.1 | 620 KB | : 100% 1.0/1 [00:02<00:00, 2.71s/it]

libgomp-14.2.0 | 449 KB | : 100% 1.0/1 [00:02<00:00, 2.78s/it]

libgomp-14.2.0 | 449 KB | : 100% 1.0/1 [00:02<00:00, 2.78s/it]

gmp-6.3.0 | 449 KB | : 100% 1.0/1 [00:03<00:00, 2.82s/it]

gmp-6.3.0 | 449 KB | : 100% 1.0/1 [00:03<00:00, 2.82s/it]

... (more hidden) ...

... (more hidden) ...

libumfpack-6.3.5 | 424 KB | : 100% 1.0/1 [00:03<00:00, 2.86s/it]

libumfpack-6.3.5 | 424 KB | : 100% 1.0/1 [00:03<00:00, 2.86s/it]

fmilib-2.4.1 | 383 KB | : 100% 1.0/1 [00:03<00:00, 2.95s/it]

fmilib-2.4.1 | 383 KB | : 100% 1.0/1 [00:03<00:00, 2.95s/it] numpy-2.2.4 | 8.6 MB | : 100% 1.0/1 [00:03<00:00, 3.25it/s]

libspqr-4.3.4 | 213 KB | : 100% 1.0/1 [00:03<00:00, 2.99s/it]

scipy-1.15.2 | 16.4 MB | : 100% 1.0/1 [00:04<00:00, 1.48it/s]

```
Preparing transaction: - 22\ 22| 22done

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

```
In [9]: # Notes of BPL_IEC_validation
```

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

- FMU BPL_IEC_Column_system_linux_om_me
- Setup-file BPL_IEC_explore

Cloning into 'BPL_IEC_validation'...

```
In [11]: %cd BPL_IEC_validation
```

 $/{\tt content/BPL_IEC_validation}$

BPL IEC validation

Author: Jan Peter Axelsson

```
In [12]: run -i BPL_IEC_explore.py
```

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

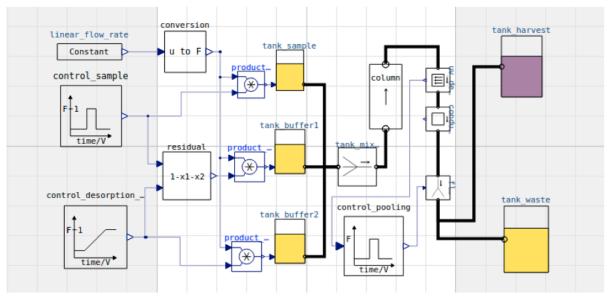
Note that both disp() and describe() takes values from the last simulation and the command process_diagram() brings up the main configuration

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

```
In [13]: plt.rcParams['figure.figsize'] = [30/2.54, 24/2.54]
```

In [14]: # The process diagram is made outside Modelica to illustrate the configuration
process_diagram()

No processDiagram.png file in the FMU, but try the file on disk.



```
In [15]: describe('chromatography'); # print(); describe('liquidphase')
```

Ion exchange chromatorgraphy controlled with varying salt-concentration. The pH is k ept constant.

Loading or adsorption

The parameter notation and values are the same as in the referred report. However the flow rate is here denoted F while q in the report. The column is diveded in n=8 sections and set at compilation time. The values are arbitrarly chosen in the report and the focus is on qualitative aspects of the model.

The simplified model describe only the column in terms of volume and does not distinguish a high column with a small diameter from a lower with larger diameter.

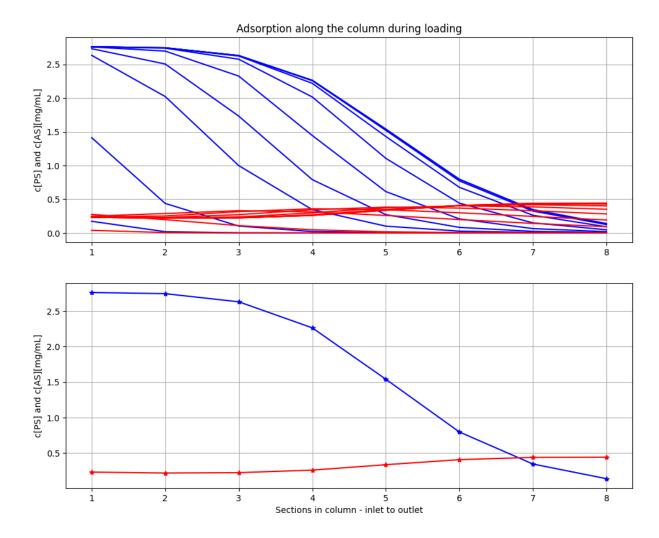
The parameters k1, k2, k3, k4 and Q_av are given relative volume and with increased column volume a larger capacity is thus obtained.

```
In [16]: # Loading of the column - try to reproduce Jonas figure 13.
         newplot(title='Adsorption along the column during loading', plotType='Loading')
         # Sample
         par(P_in=1.0, A_in=1.0, E_in=0)
         # Column properties
         par(k1=0.3, k2=0.05, k3=0.05, k4=0.3, Q_av=3.0)
         par(height=20, diameter=0.714)
         par(x_m=0.3)
         # Operation
         par(E_in_desorption_buffer=8)
         par(LFR=12)
         #par(scale volume=False)
         par(start_adsorption=0, stop_adsorption=50)
         par(start_desorption=150, stationary_desorption=450)
         par(start_pooling=220, stop_pooling=450)
         # Simulation
         simu(100)
        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)

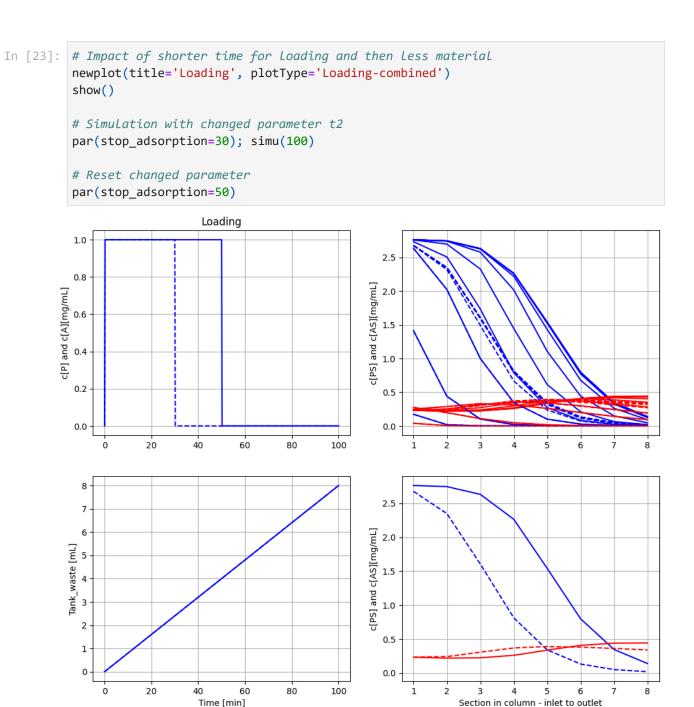
Could not find ODEPACK functions.

Could not find RADAR5
Could not find GLIMDA.



The results are the same af Figure 13 in [1].

```
# We just check that we had the same volume flow rate as Jonas
In [17]:
         describe('F')
         : 0.08
In [18]: describe('V')
        Column volume total - derived : 8.008 [ mL ]
In [19]:
         model.get('column.x_m')
Out[19]: array([0.3])
In [20]:
         model.get('column.V_m')
Out[20]:
         array([2.40235705])
         #describe('column.n')
In [21]:
         model.get('column.column_section[1].V_m')
In [22]:
Out[22]: array([0.30029463])
```



To the left the inlet loading over time. To the right upper concentration along the column at different times and in steady states finally To the right lower concentrations along the column in steady state.

We see that a shorter time and then less material makes less of the column capacity used.

Note that the flow through the column is constant despite change from sample to just buffer 1, and shown in how the volume of the waste tank increase with time.

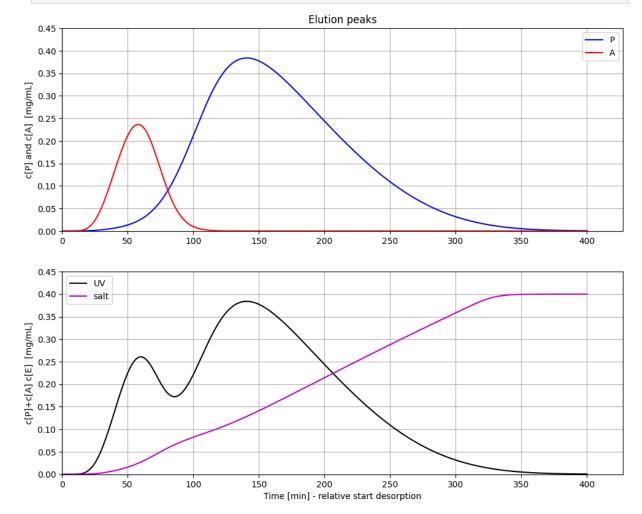
Elution or desorption

```
In [24]: # Elution of the column
    newplot(title='Elution peaks', plotType='Elution')

# Sample
    par(P_in=1, A_in=1.0, E_in=0)

# Operation
    par(E_in_desorption_buffer=8)
    par(LFR=12.0, start_adsorption=0, stop_adsorption=50, start_desorption=150, station

# Simulation
    simu(550)
```



The results are the same af Figure 14 in [1].

The upper diagrams shows the column outlet concentrations of P and A over time.

The lower diagram shows the sum (or possibly the UV signal) at column outlet as well as the salt concentration. We have some separation between the two peaks.

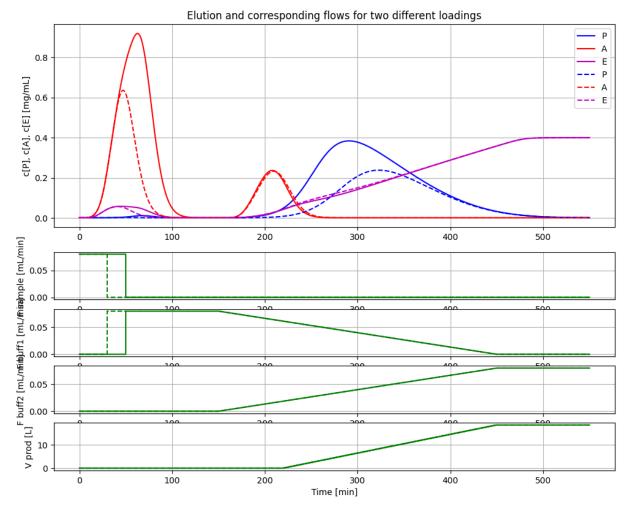
Note that the salt concentration deviates slightly from the linear increse between time 50 to 100. This is due to ion interaction with P and A in the column. This is phenomenon can also be seen in real data. The ion-salt concentration is scaled with factor 0.05 to get comparable concentrations to P and A.

I have here simulated time 150 of adsorbtion and then started elution. Here is time counted as zero at time of start of elutions. Not sure how long Jonas simulated to get steady state before he did elution.

```
In [25]: # More complete visualization of the elution phase and the different flows
    newplot(title='Elution and corresponding flows for two different loadings', plotTyp
    par(stop_adsorption=50); simu(550)

# Simulation with changed parameter t2
    par(stop_adsorption=30); simu(550)

# Reset changed parameter
    par(stop_adsorption=50)
```

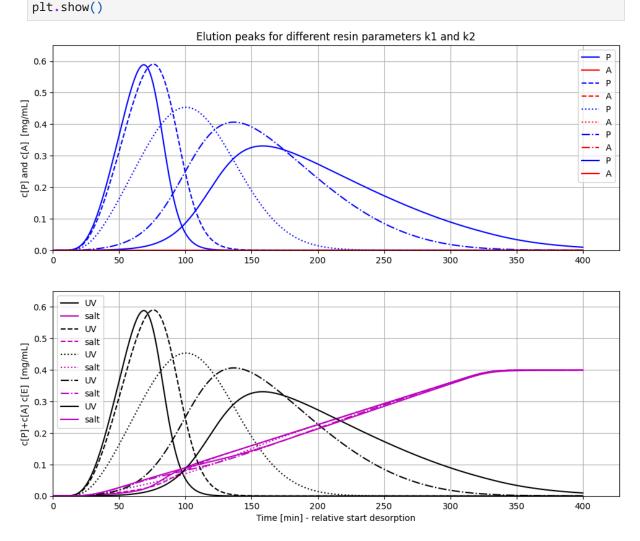


Here a diagram that shows the peaks at the outlet as shown in the previous diagram. Below the flow rates of the three different sorces. Here time is 0 at start of adsorbtion and elution starts at time 150.

Automatic pooling based on UV-mmeasurement is tested in another notebook.

Change of resin properties

```
In [26]:
         # Elution of the column
         newplot(title='Elution peaks for different resin parameters k1 and k2', plotType='E
         # Sample
         par(P_in=1, A_in=0.0, E_in=0)
         # Operation
         par(E_in_desorption_buffer=8)
         par(LFR=12.0, start_adsorption=0, stop_adsorption=50, start_desorption=150, station
         # Simulations
         par(k1=0.05, k2=0.50); simu(550)
         par(k1=0.05, k2=0.25); simu(550)
         par(k1=0.05, k2=0.05); simu(550)
         par(k1=0.25, k2=0.05); simu(550)
         par(k1=0.50, k2=0.05); simu(550)
         # Adjust diagrams
         ax1.set_ylim([0, 0.65])
         ax2.set_ylim([0, 0.65])
```



The results are the same af Figure 17 in [1].

Summary

Three important diagrams Figure 13, 14 and 17 in the original report [1] were reproduced and the implementation in Modelica used here is considered validated.

The model is now extended with an improved parametrization of the column that match the industrial practice.

Acknowledgement

The author thank Karl Johan Brink for sharing his know-how of chromatography operation. He has especially given input to how to parametrize the model in terms often used in the industrial practice.

References

- 1. Månsson, Jonas, "Control of chromatography comlumn in production scale", Master thesis TFRT-5599, Department of Automatic Control, LTH, Lund Sweden, 1998.
- 2. Pharmacia LKB Biotechnology. "Ion Exchange chromatography. Principles and Mathods.", 3rd edition, 1991.

Appendix

```
In [27]: describe('MSL')
        MSL: 3.2.3 - used components: RealInput, RealOutput, CombiTimeTable, Types
In [28]: system_info()
        System information
         -OS: Linux
         -Python: 3.11.11
         -Scipy: not installed in the notebook
         -PyFMI: 2.16.3
         -FMU by: OpenModelica Compiler OpenModelica 1.25.0~dev-133-ga5470be
         -FMI: 2.0
         -Type: FMUModelME2
         -Name: Column_system
         -Generated: 2024-11-07T11:26:29Z
         -MSL: 3.2.3
         -Description: Bioprocess Library version 2.3.0
         -Interaction: FMU-explore version 1.0.0
In [28]:
```