BPL_TEST2_Fedbatch - demo

This notebook just produce the Figure 2 in the paper "Design ideas behind Bioprocess Library for Modelica", by J P Axelsson, presented in the 15th International Modelica Conference in Aachen, Germany, October 9-11, 2023.

Unfortunately the Linux FMU from OpenModelica does not work and therefore Windows FMU from JModelica is used. Thus Google Colab cannot run the example.

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

- describe() - describe culture, broth, parameters, variables with values/units

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

```
In [2]: %matplotlib inline
plt.rcParams['figure.figsize'] = [25/2.54, 25/2.54]

In [3]: process_diagram()

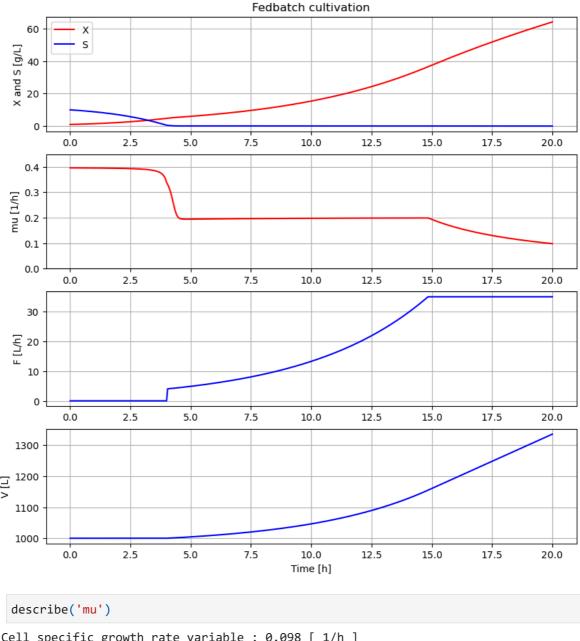
dosagescheme

feedtank

bioreactor

feedtank
```

```
In [4]: # Simulation with default values of the process
newplot(plotType='TimeSeries')
ax2.set_ylim([0, 0.45])
init(V_0=1000, VX_0=1*1e3, VS_0=10*1e3)
par(feedtank_S_in=600, Ks=0.1, Y=0.40)
par(t_start=4, F_start=4.0, mu_feed=0.2, F_max=35)
simu(20)
```



```
In [5]: describe('mu')
    Cell specific growth rate variable : 0.098 [ 1/h ]
In [6]: describe('parts')
    ['bioreactor', 'bioreactor.culture', 'dosagescheme', 'feedtank', 'liquidphase', 'MSL']
In [7]: describe('MSL')
    MSL: RealInput, RealOutput
In [8]: system_info()
```

 ${\tt System \ information}$

-OS: Windows
-Python: 3.12.9

-Scipy: not installed in the notebook

-PyFMI: 2.16.3

-FMU by: JModelica.org

-FMI: 2.0

-Type: FMUModelCS2

-Name: BPL_TEST2.Fedbatch
-Generated: 2023-03-30T09:13:00

-MSL: 3.2.2 build 3

-Description: Bioprocess Library version 2.1.1

-Interaction: FMU-explore version 0.9.8

In []: