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.

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In [1]: run -i BPL_TEST2_Fedbatch_explore.py
        Windows - run FMU pre-compiled JModelica 2.14
        Model for bioreactor has been setup. Key commands:
         - par()

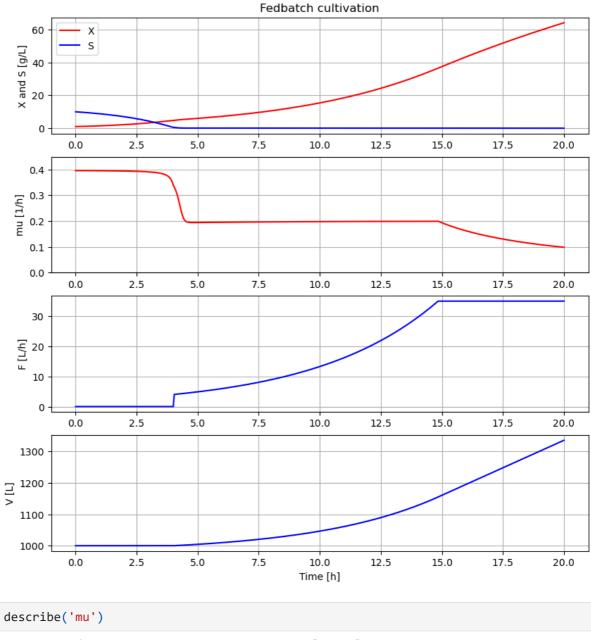
    change of parameters and initial values

    change initial values only

         - init()

    simulate and plot

         - simu()
         - newplot() - make a new plot
                     - show plot from previous simulation
         - show()
         - disp() - display parameters and initial values from the last simulation
         - describe() - describe culture, broth, parameters, variables with values / unit
        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()
In [2]: %matplotlib inline
        plt.rcParams['figure.figsize'] = [25/2.54, 25/2.54]
In [3]: # 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)
```



System information

-OS: Windows
-Python: 3.10.6

-Scipy: not installed in the notebook

-PyFMI: 2.10.3

-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 0.9.7

In []: