BPL_TEST2_Chemostat - demo

In [1]: run -i BPL_TEST2_Chemostat_fmpy_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()show plot from previous simulationdisp()display parameters and initial values from the last simulation
- describe() describe culture, broth, parameters, variables with values/uni

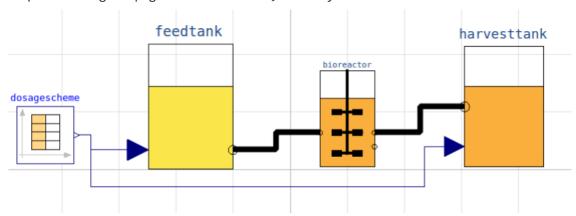
Note that both disp() and describe() takes values from the last simulation and the command process_diagram() brings up the main configuration

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

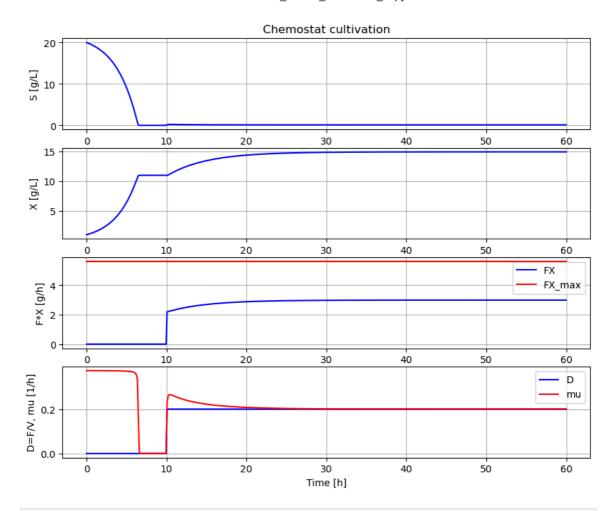
```
In [2]: %matplotlib inline
        plt.rcParams['figure.figsize'] = [25/2.54, 20/2.54]
```

In [3]: process_diagram()

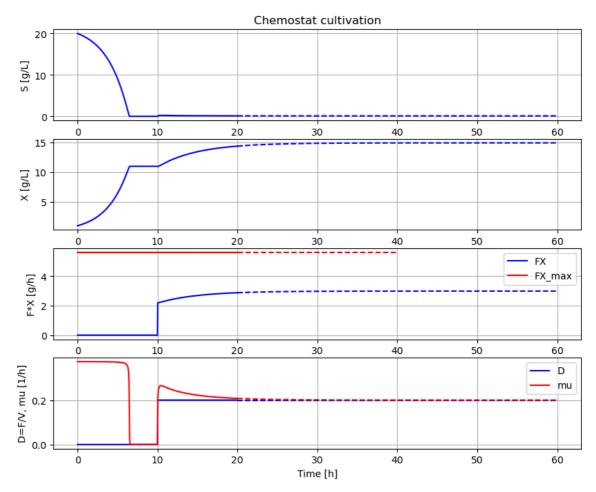
No processDiagram.png file in the FMU, but try the file on disk.



```
In [4]: newplot()
        par(Y=0.50, qSmax=0.75, Ks=0.1)
                                                      # Culture parameters
        init(V start=1.0, VX start=1.0, VS start=20) # Bioreactor startup
        par(S_in=30, t0=0, F0=0, t1=10, F1=0.2)
                                                      # Substrate feeding
        simu(60)
```



```
In [5]: # Check simu('cont')
    newplot()
    simu(20)
    simu(40,'cont')
```



System information

-OS: Windows
-Python: 3.9.16

-Scipy: not installed in the notebook

-FMPy: 0.3.15

-FMU by: JModelica.org

-FMI: 2.0 -Type: CS

-Name: BPL_TEST2.Chemostat -Generated: 2024-05-12T20:53:47

-MSL: 3.2.2 build 3

-Description: Bioprocess Library version 2.2.0 -Interaction: FMU-explore for FMPy version 1.0.0

In []: