## BPL\_TEST2\_Chemostat - demo

In [1]: run -i BPL TEST2 Chemostat fmpy explore.py

Linux - run FMU pre-compiled OpenModelica

Model for the process has been setup. Key commands:

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

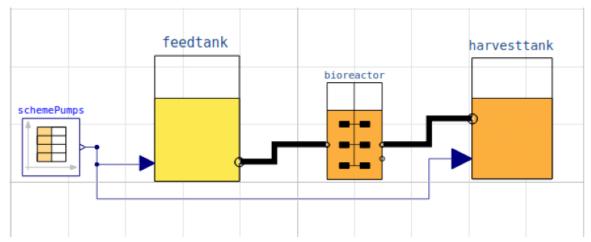
Note that both disp() and describe() takes values from the last simulation and the command process\_diagram() brings up the main configuration

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

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

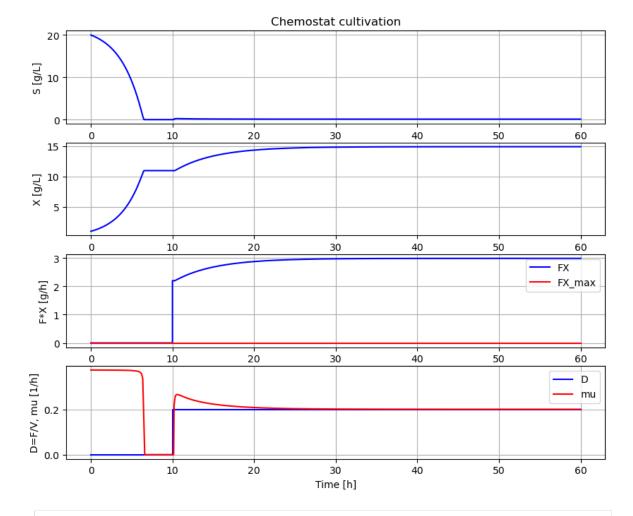
In [3]: process\_diagram()

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



```
In [4]: newplot()
     par(Y=0.50, gSmax=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)
```

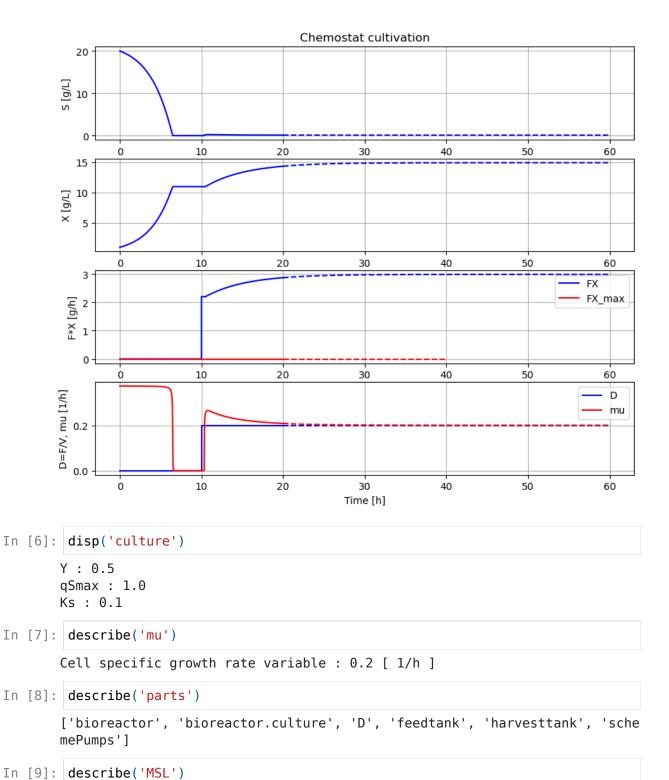
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```
In [5]: # Check simu('cont')
 newplot()
 simu(20)
 simu(40,'cont')
```

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In [10]: system\_info()



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MSL: 4.1.0 - used components: RealInput, RealOutput, CombiTimeTable, Types

## System information

-OS: Linux

-Python: 3.12.11

-Scipy: not installed in the notebook

-FMPy: 0.3.26

-FMU by: OpenModelica Compiler OpenModelica 1.26.0~dev-200-gcb3254b

-FMI: 2.0 -Type: ME

-Name: BPL.Examples\_TEST2.Chemostat -Generated: 2025-07-28T07:59:17Z

-MSL: 4.1.0

-Description: Bioprocess Library version 2.3.1 -Interaction: FMU-explore for FMPy version 1.0.1

## In [11]: !lsb\_release -a

No LSB modules are available.

Distributor ID: Ubuntu

Description: Ubuntu 24.04.3 LTS

Release: 24.04 Codename: noble

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