## 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
- 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

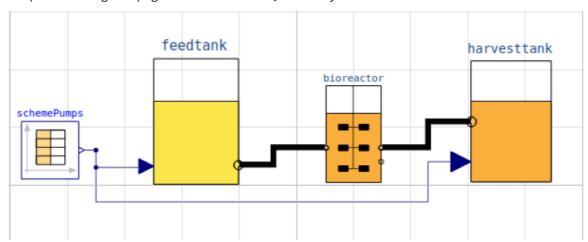
Note that both disp() and describe() takes values from the last simulation and the command process\_diagram() brings up the main configuration

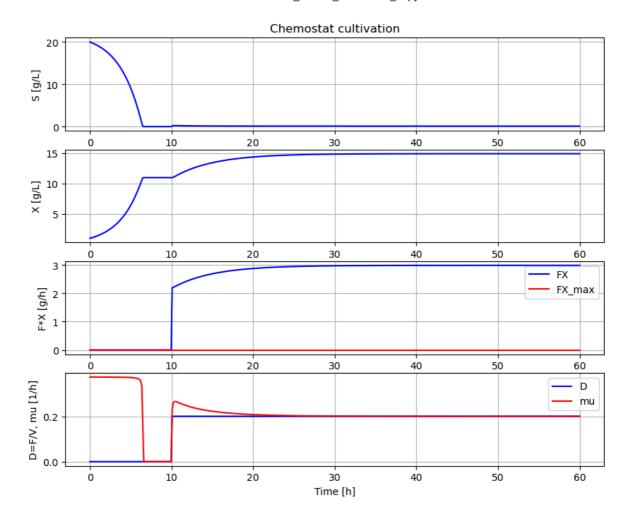
Brief information about a command by help(), eg help(simu)
Key system information is listed with the command system\_info()

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

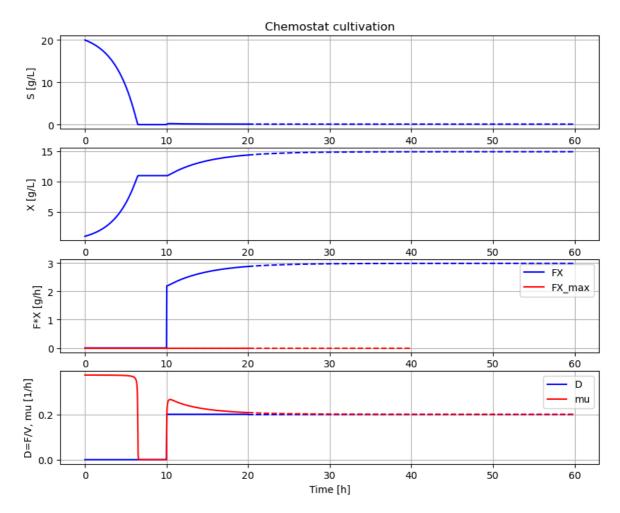
In [3]: process\_diagram()

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





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



```
In [6]: disp('culture')

Y: 0.5
qSmax: 1.0
Ks: 0.1

In [7]: describe('mu')
Cell specific growth rate variable: 0.2 [ 1/h ]

In [8]: describe('parts')
['bioreactor', 'bioreactor.culture', 'D', 'feedtank', 'harvesttank', 'MSL', 'sche mePumps']

In [9]: describe('MSL')
MSL: RealInput, RealOutput, CombiTimeTable, Types

In [10]: system_info()
```

System information -OS: Windows -Python: 3.12.3

-Scipy: not installed in the notebook

-FMPy: 0.3.20

-FMU by: JModelica.org

-FMI: 2.0 -Type: CS

-Name: BPL.Examples\_TEST2.Chemostat -Generated: 2024-10-24T08:57:28

-MSL: 3.2.2 build 3

-Description: Bioprocess Library version 2.2.2 - GUI -Interaction: FMU-explore for FMPy version 1.0.1

In [ ]: