## BPL\_TEST2\_Chemostat - demo

In [1]: run -i BPL\_TEST2\_Chemostat\_explore.py

Windows - run FMU pre-compiled JModelica 2.14

Model for the process 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
- $\operatorname{disp}()$   $\operatorname{display}$  parameters and initial values from the last simulation
- describe() describe culture, broth, parameters, variables with values/units

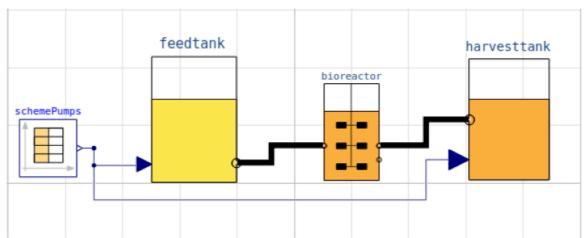
Note that both disp() and describe() takes values from the last simulation and the command process\_diagram() brings up the main configuration

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

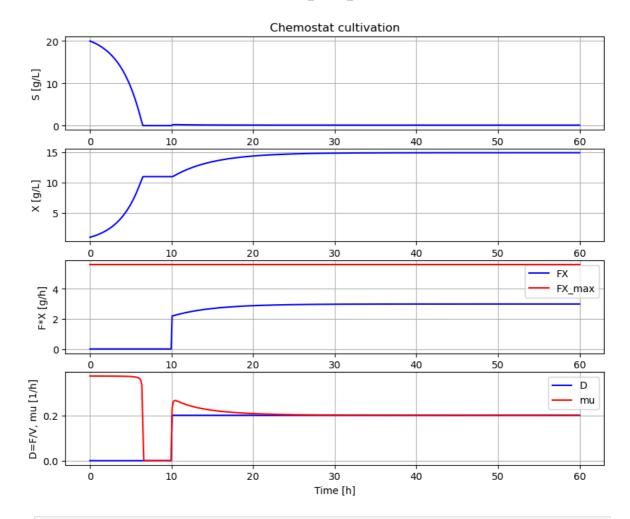
```
In [2]: %matplotlib inline
  plt.rcParams['figure.figsize'] = [25/2.54, 20/2.54]
```

In [3]: process\_diagram()

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



```
In [4]: #fmu_model ='xBPL_TEST2_Chemostat_linux_om_me.fmu'
#model = load_fmu(fmu_model, log_level=0)
```



In [6]: # The maximal biomass productivity FX\_max [g/h] marked red in the diagram above
# can be calculated for CSTR from the FMU and is
cstrProdMax(model)

```
Out[6]: np.float64(5.625)
```

```
In [7]: describe('cstrProdMax')
```

Calculate from the model maximal chemostat productivity FX\_max : 5.625 [ g/h ]

```
In [8]: disp('culture')
```

Y : 0.5 qSmax : 0.75 Ks : 0.1

In [9]: describe('mu')

Cell specific growth rate variable : 0.2 [ 1/h ]

```
In [10]: describe('parts')
```

['bioreactor', 'bioreactor.culture', 'D', 'feedtank', 'harvesttank', 'MSL', 'sche mePumps']

In [11]: describe('MSL')

MSL: RealInput, RealOutput, CombiTimeTable, Types

In [12]: system\_info()

 ${\tt System \ information}$ 

-OS: Windows
-Python: 3.12.11

-Scipy: not installed in the notebook

-PyFMI: 2.17.4

-FMU by: JModelica.org

-FMI: 2.0

-Type: FMUModelCS2

-Name: BPL.Examples\_TEST2.Chemostat -Generated: 2025-07-26T09:39:44

-MSL: 3.2.2 build 3

-Description: Bioprocess Library version 2.3.1

-Interaction: FMU-explore version 1.0.0

In [ ]: