

BPL_TEST2_Chemostat - demo

In [1]: `run -i BPL_TEST2_Chemostat_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

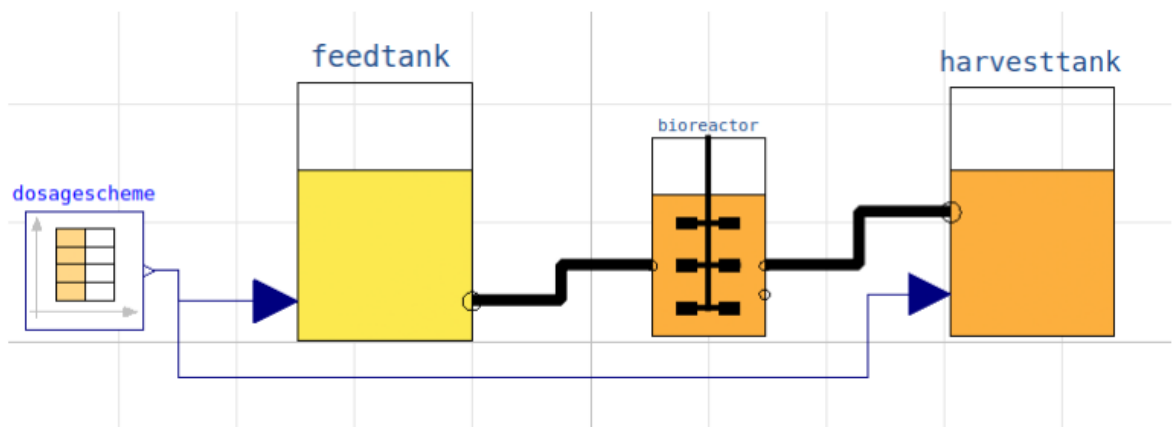
Note that both `disp()` and `describe()` takes values from the last simulation and the command `process_diagram()` brings up the main configuration

Brief information about a command by `help()`, eg `help(simu)`

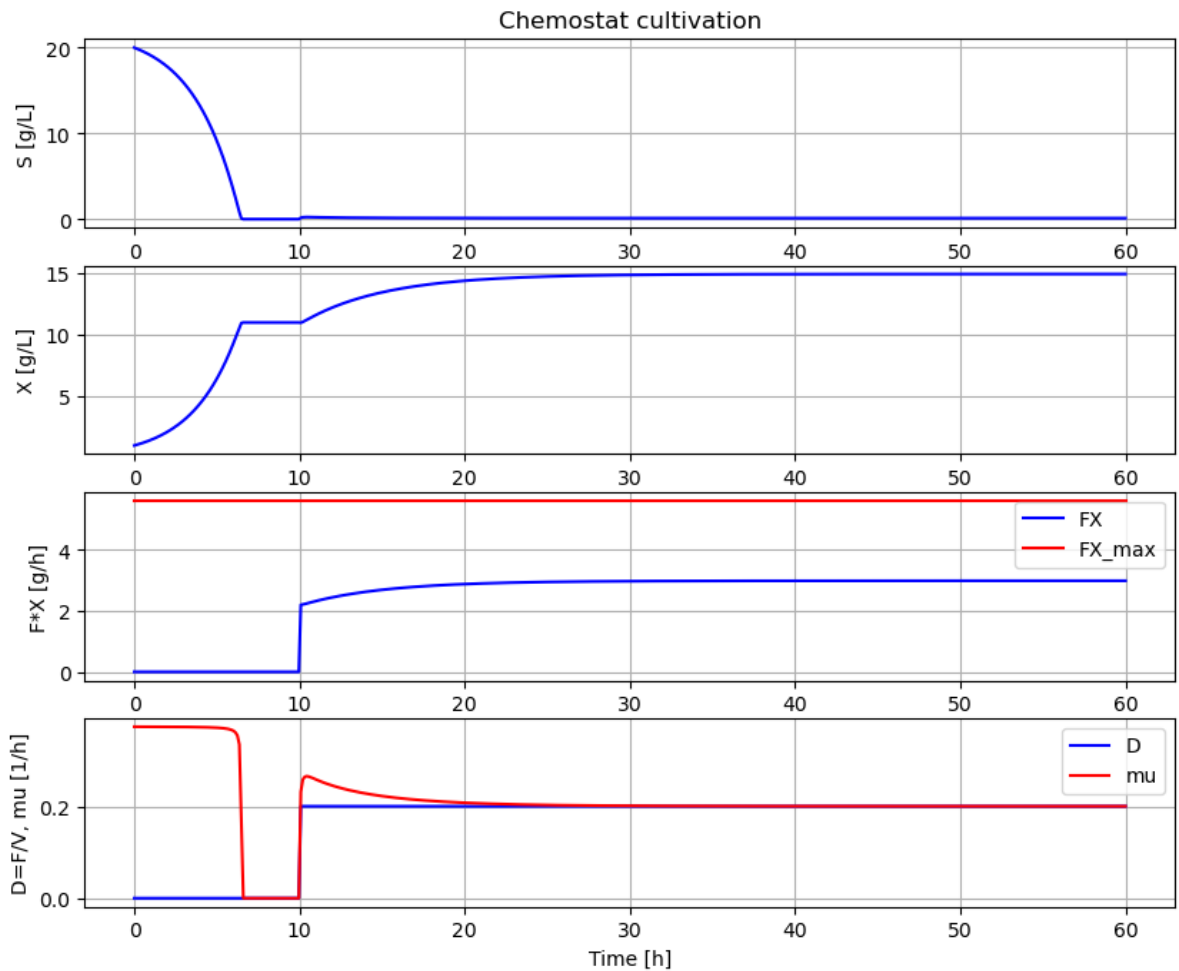
Key system information is listed with the command `system_info()`

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

In [3]: `process_diagram()`



In [4]: `newplot()`
`par(Y=0.50, qSmax=0.75, Ks=0.1) # Culture parameters`
`init(V_0=1.0, VX_0=1.0, VS_0=20) # Bioreactor startup`
`par(S_in=30, t0=0, F0=0, t1=10, F1=0.2) # Substrate feeding`
`simu(60)`



```
In [5]: # 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[5]: 5.625
```

```
In [6]: describe('cstrProdMax')
```

Calculate from the model maximal chemostat productivity FX_{max} : 5.625 [g/h]

```
In [7]: disp('culture')
```

```
Y : 0.5
qSmax : 0.75
Ks : 0.1
```

```
In [8]: describe('mu')
```

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

```
In [9]: describe('parts')
```

```
['bioreactor', 'bioreactor.culture', 'D', 'dosagescheme', 'feedtank', 'harvesttan
k', 'liquidphase', 'MSL']
```

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

MSL: RealInput, RealOutput, CombiTimeTable, Types

```
In [11]: system_info()
```

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.Chemostat
- Generated: 2023-09-02T07:45:11
- MSL: 3.2.2 build 3
- Description: Bioprocess Library version 2.1.2 prel
- Interaction: FMU-explore version 0.9.8

In []: