

BPL_TEST2_Chemostat - demo

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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

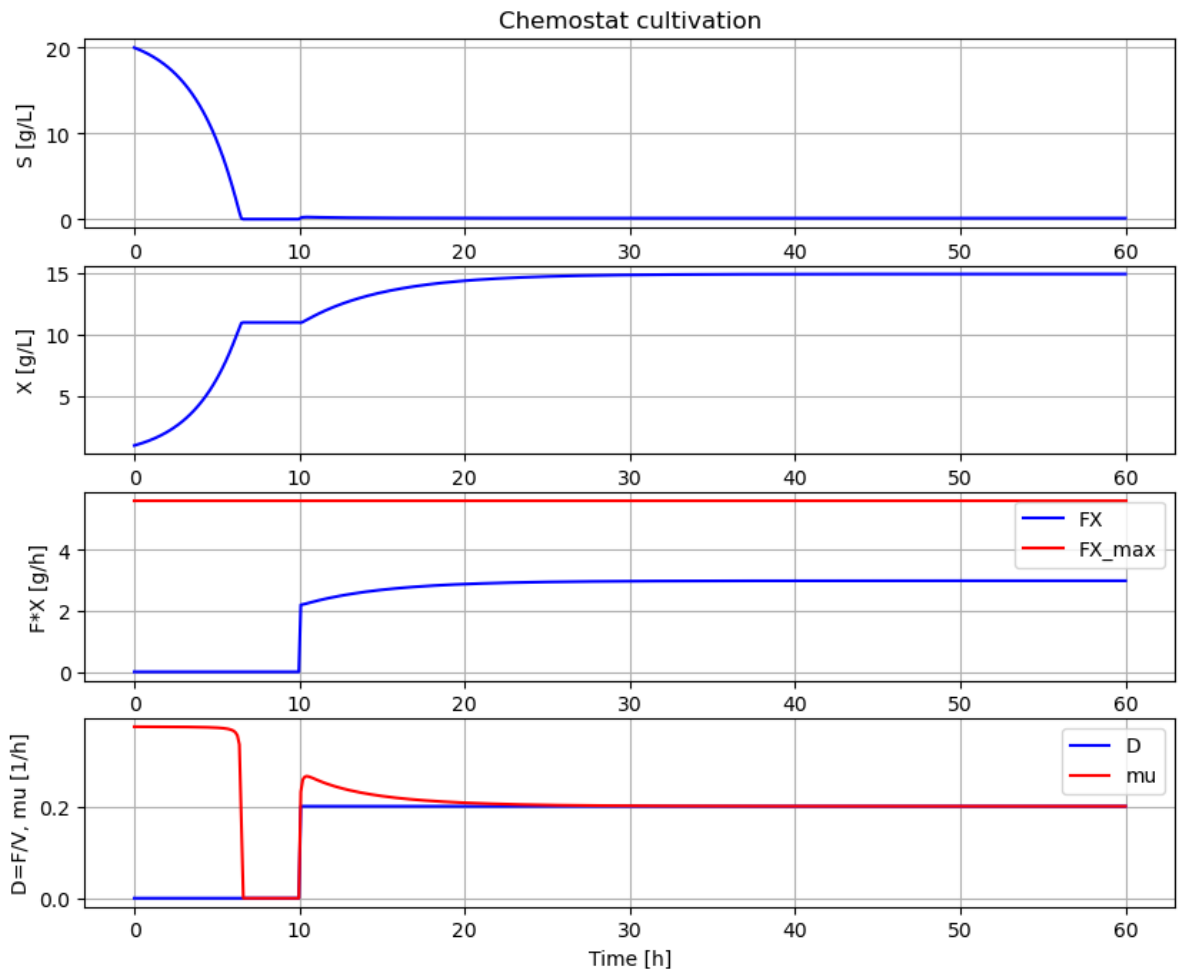
Note that both disp() and describe() takes values from the last simulation

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]: 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 [4]: # 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[4]: 5.625
```

```
In [5]: describe('cstrProdMax')
```

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

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

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

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

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

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

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

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

MSL: 3.2.2 build 3 - used components: RealInput, RealOutput, CombiTimeTable, Types

```
In [10]: system_info()
```

System information

- OS: Windows
- Python: 3.10.6
- Scipy: not installed in the notebook
- PyFMI: 2.9.8
- FMU by: JModelica.org
- FMI: 2.0
- Type: FMUModelCS2
- Name: BPL_TEST2.Chemostat
- Generated: 2022-10-17T19:59:47
- MSL: 3.2.2 build 3
- Description: Bioprocess Library version 2.1.0
- Interaction: FMU-explore version 0.9.5

In []: