

```
disp('culture')
```

```
⇒ Y : 0.5  
   qSmax : 1.0  
   Ks : 0.1
```

```
describe('mu')
```

```
⇒ Cell specific growth rate variable : 0.12 [ 1/h ]
```

```
describe('parts')
```

```
⇒ ['bioreactor', 'bioreactor.culture', 'dosagescheme', 'feedtank']
```

```
describe('MSL')
```

```
⇒ MSL: 3.2.3 – used components: RealInput, RealOutput
```

```
system_info()
```

```
⇒  
System information  
-OS: Linux  
-Python: 3.10.12  
-Scipy: not installed in the notebook  
-PyFMI: 2.14.0  
-FMU by: OpenModelica Compiler OpenModelica 1.23.1  
-FMI: 2.0  
-Type: FMUModelME2  
-Name: BPL.Examples_TEST2.Fedbatch  
-Generated: 2024-08-31T07:47:59Z  
-MSL: 3.2.3  
-Description: Bioprocess Library version 2.2.1 – GUI  
-Interaction: FMU-explore version 1.0.0
```

Start coding or [generate](#) with AI.