

Pure Modelica Unit Testing: From Mathematical Algorithms to Physical Modeling

OpenModelica Annual Workshop 2019

Atiyah Elsheikh
Mathemodica.com

4th February 2019

Atiyah.Elsheikh@mathemodica.com

Outline

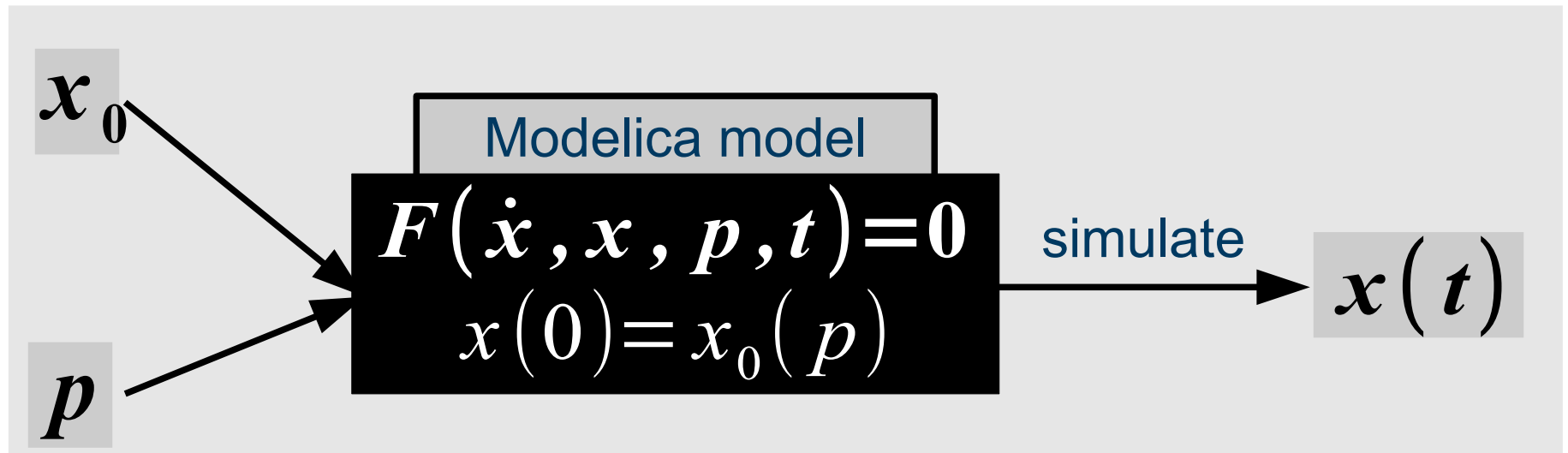
- First testing experiences
 - Mathematical algorithms in Modelica
- Unit testing in Modelica
 - existing technologies
- A bit more “formal” Unit testing
 - Upgrading to Physical Modeling

Motivation

Wiechert, W., Noack, S., and Elsheikh, A. (2010).
Modeling languages for biochemical network
simulation: Reaction vs equation based approaches.
Advances in Biochemical Engineering / Biotechnology

“Simulation tools not only perform numerical
solutions based on the system equations but
also assist the modeler in systems analysis.
Doubtlessly the most important systems analysis
tool is Sensitivity Analysis ...

SA of Modelica Models

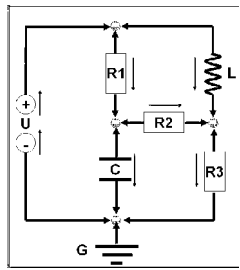


Dynamic Parameter Sensitivities (DPS)

$$\frac{\partial x}{\partial p}(t)$$

$$\frac{\partial x}{\partial x_0}(t)$$

ADModelica (2007)



OMC
Compiler

$$\begin{aligned}
 R1.v &= R1.R \cdot R1.i \\
 R2.v &= R2.R \cdot R2.i \quad Lv = LL \cdot Li \\
 R3.v &= R3.R \cdot R3.i \\
 U.v &= -U_0 \quad \dot{C}.v = C.i / C.C \\
 G.in.v &= 0
 \end{aligned}$$

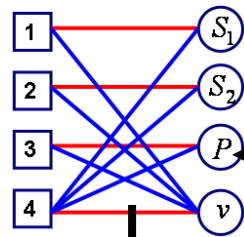
XML
Modelica

```

- <equation scolumn="27" slime="51">
- <equ_equality scolumn="27" slime="51">
- <add operation="binary" scolumn="18" slime="51">
- <add operation="binary" scolumn="9" slime="51">
  <component_reference ident="L_n_i" scolumn="3" slime="51">
  <component_reference ident="R1_p_i" scolumn="11" slime="51">
  </add>
  <component_reference ident="R2_p_i" scolumn="20" slime="51">
  </add>
  <real_literal scolumn="29" slime="51" value="0.0"/>
</equ_equality>
</equation>

```

ADModelica



XML
Parser

Visualizer

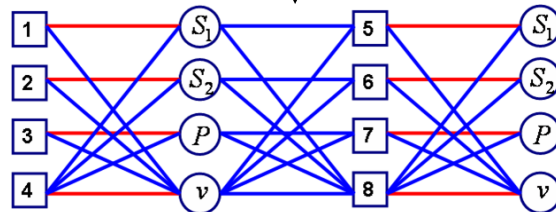
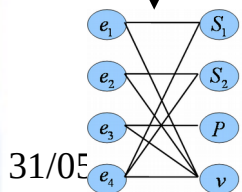
Analyzer

Unparser

```

equation
(R1_R+R1_p_i) = (L_n_v-C_p_v);
for ad_i in 1:G_N loop
  ((R1_p_i+g_R1_R[ad_i])+(R1_R+g_R1_p_i(ad_i))=
  (g_L_n_v[ad_i]-g_C_p_v[ad_i]);
end for;

```



Sample of generated code (I/II)

- Declaration part

```
model ADSimpleReaction
  Real S(start=1);
  Real P(start=0);
  Real v;
  Real[3] g_S, g_P, g_v;
  parameter Real vmax=1;
  constant Real[3] g_vmax={1,0,0};
  parameter Real k=1;
  constant Real[3] g_k={0,1,0};
  parameter Real Ik=1;
  constant Real[3] g_Ik={0,0,1};
protected
  Real loc01, loc02, loc03, loc04, loc05;
  Real g loc01, g loc02, g loc03, g loc04, g loc05;
```

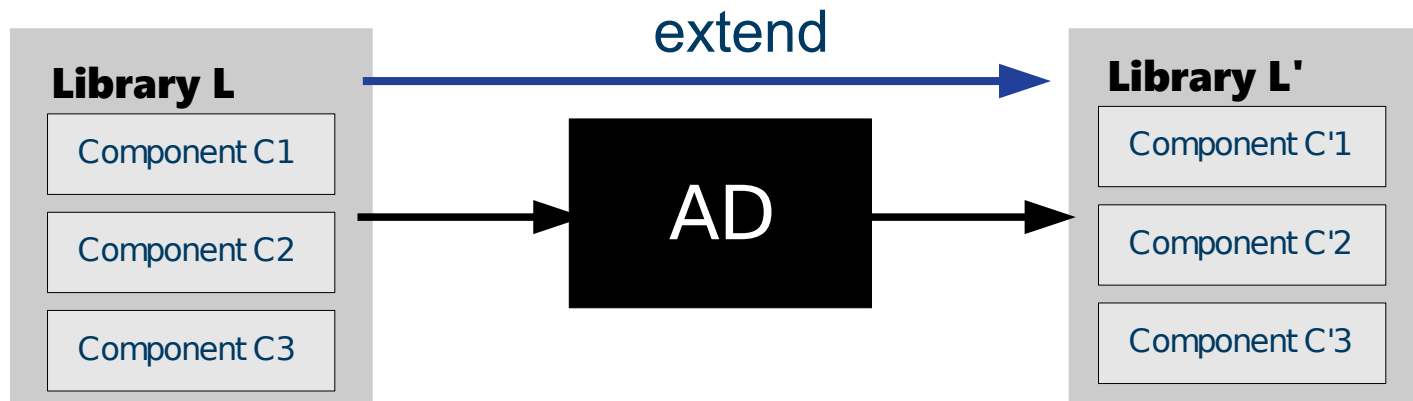
Sample of generated code

(II/II)

- Equation part

```
equation
  der(S)=-v;
  der(P)=v;
  //v = vmax * (S/(S+k)) * (Ik/(P+Ik));
algorithm
  loc01 := vmax*S;
  loc02 := S+k;
  loc03 := loc01/loc02;
  loc04 := P+Ik;
  loc05 := Ik/loc04;
  v      := loc03*loc05;
//Derivatives:
equation
  for i in 1:3 loop
    der(g_S[i])=-g_v[i];
    der(g_P[i])=g_v[i];
  end for;
algorithm
  for i in 1:3 loop
    g_loc01 := g_vmax[i]*S+vmax*g_S[i];
    g_loc02 := g_S[i]+g_k[i];
    g_loc03 := (g_loc01*loc02-loc01*g_loc02)/(loc02^2);
    g_loc04 := g_P[i]+g_Ik[i];
    g_loc05 := (g_Ik[i]*loc04-Ik*g_loc04)/(loc04^2);
    g_v[i]  := g_loc03*loc05+loc03*g_loc05;
  end for;
end ADSimpleReaction;
```

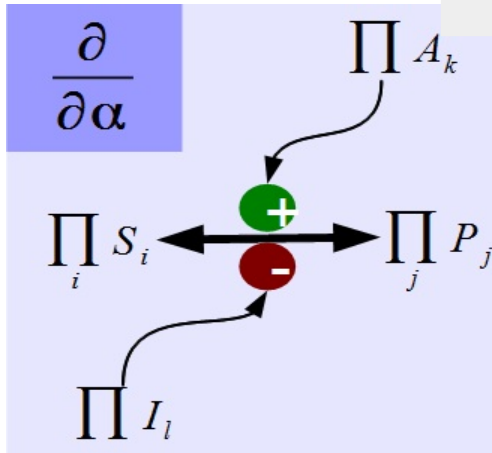
Algorithmic Differentiation (AD) of Modelica libraries (2011)



Algorithmic differentiation Modelica Libraries

The ADGenKinetics Library

2011

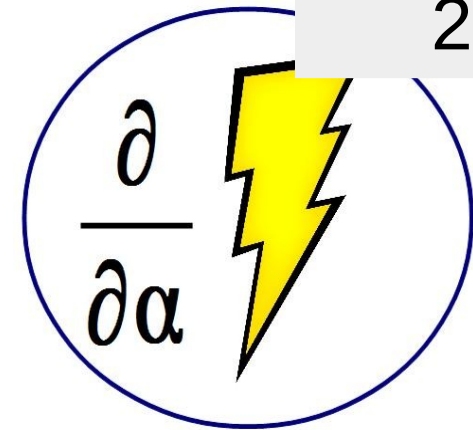


Algorithmically differentiated Modelica library for biochemical reaction networks

<https://github.com/modelica-3rdparty/ADGenKinetics>

The ADMSL Library

2014



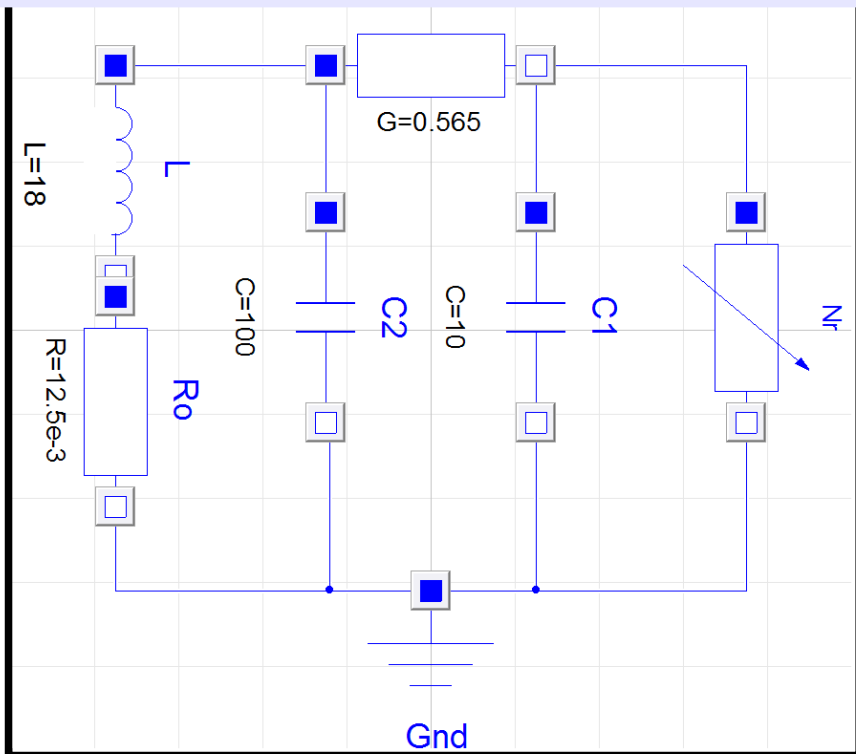
Serves as an example of algorithmically differentiated Modelica library

<https://github.com/AIT-CES-LAB/ADMSL>

Chua Circuit

Standard Modelica model for an electrical circuit

`Modelica.Electrical.Analog.Basic.Examples.ChuaCircuit`

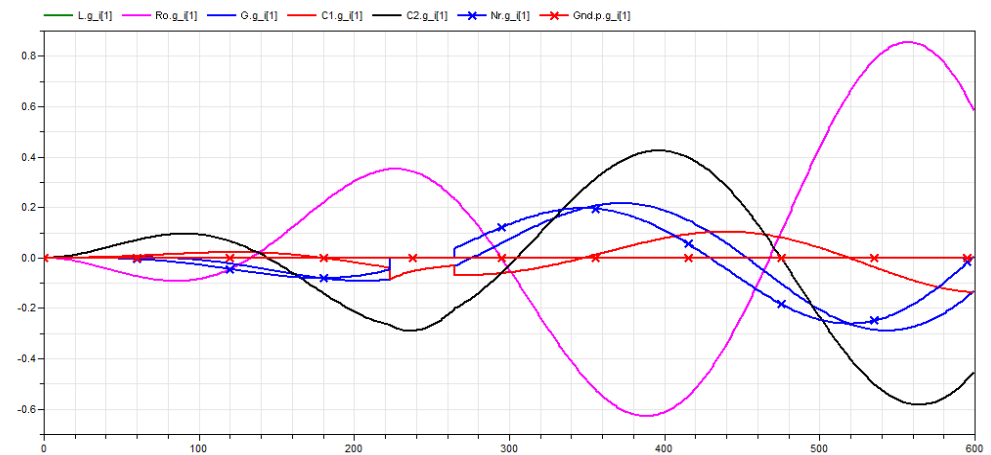
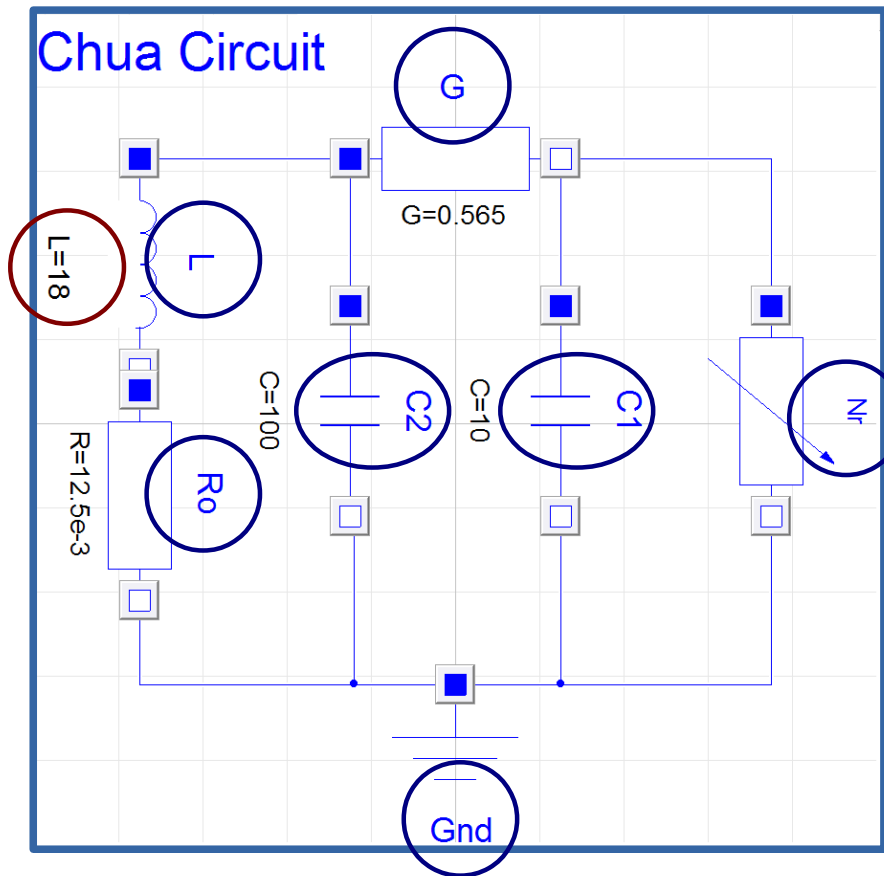


This examples simulates the current and voltage at all components

Chua Circuit importing ADMSL

The sensitivities of the current w.r.t. L

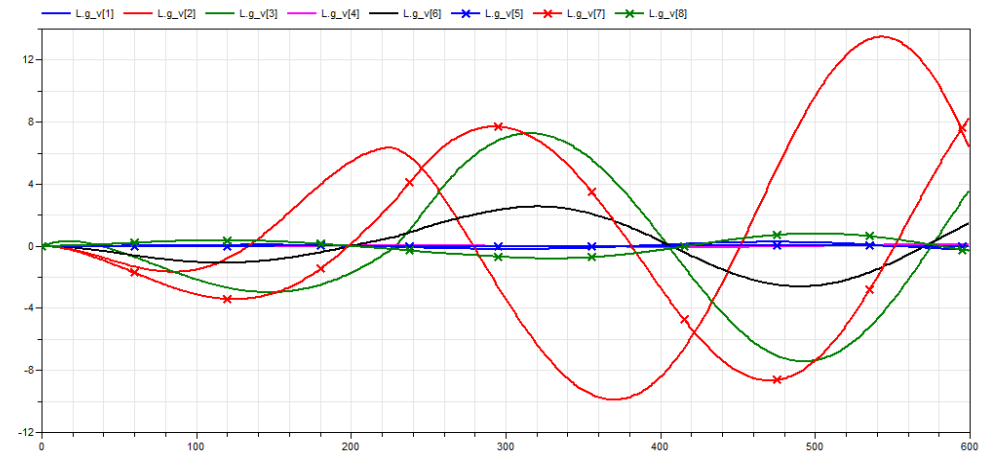
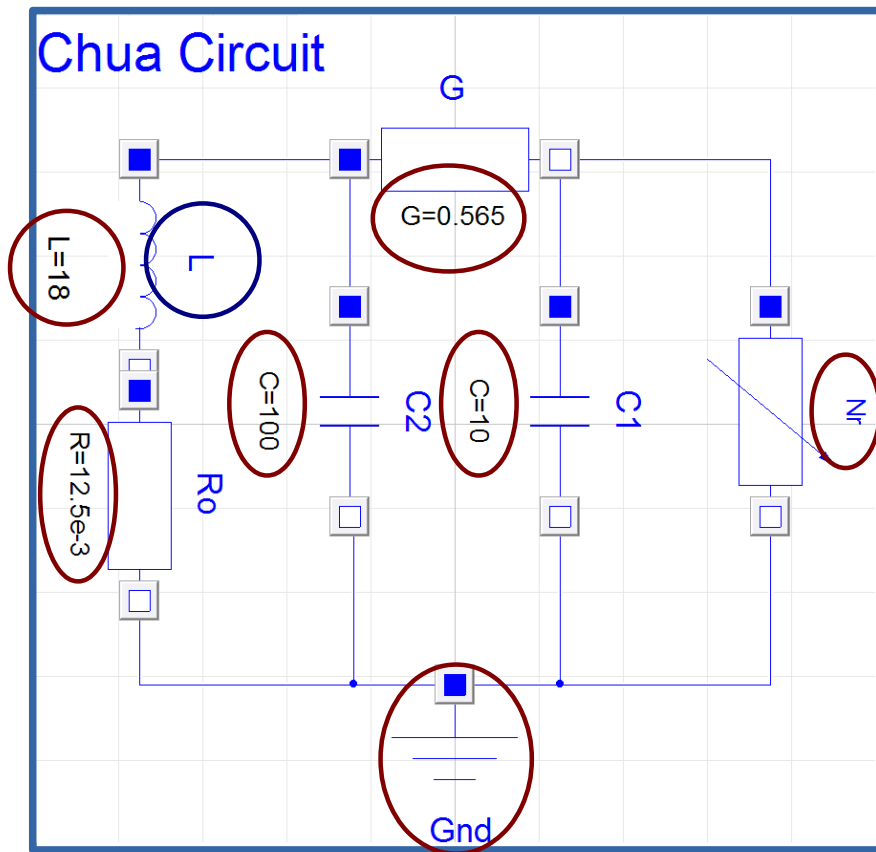
$$\frac{\partial X.i}{\partial L.L}$$



Chua Circuit importing ADMSL

The sensitivities of the voltage at L
w.r.t. all parameters

$$\frac{\partial V_L}{\partial X.p}$$



Applications of DPS

1) Modeling-Oriented

Control Coefficients, Local SA, Parameter Sweeping Studies, Model Simplification, ...

2) Statistical

Regression Analysis, Global SA, Identifiability Analysis, ...

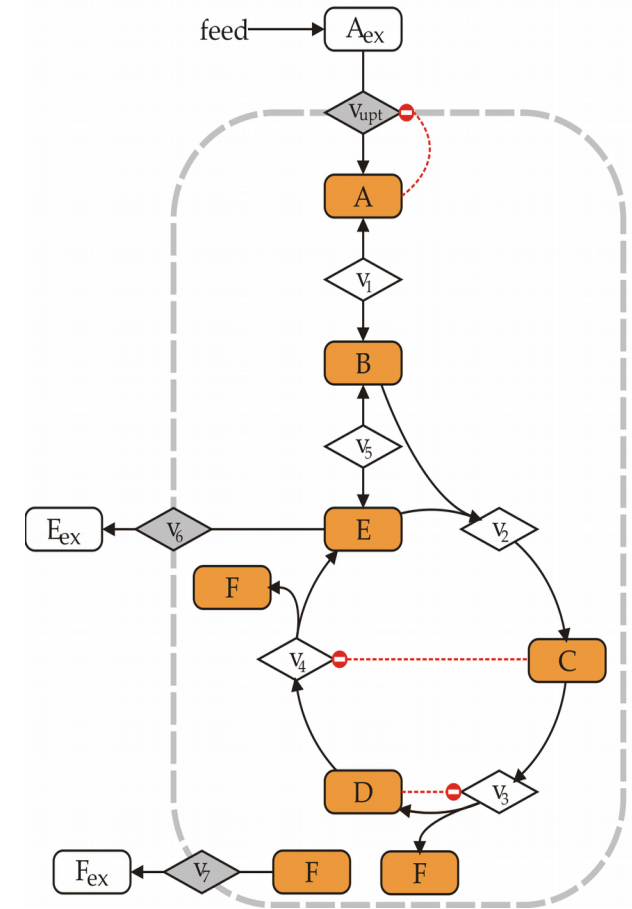
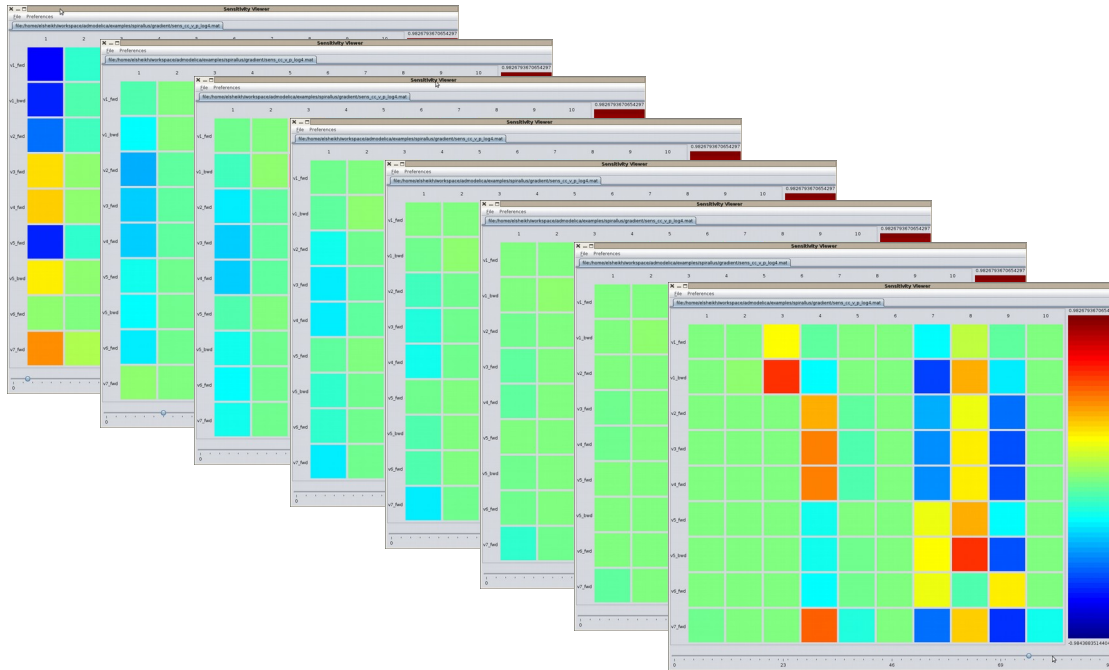
3) Optimization

Cost-functions expressed in terms of DPS

Atiyah Elsheikh and Sergei Kucherenko. (2019)
Dynamic parameter sensitivities: Summary of applications - version 1.0.
Technical Report, to appear

Scaled Parameter Sensitivities

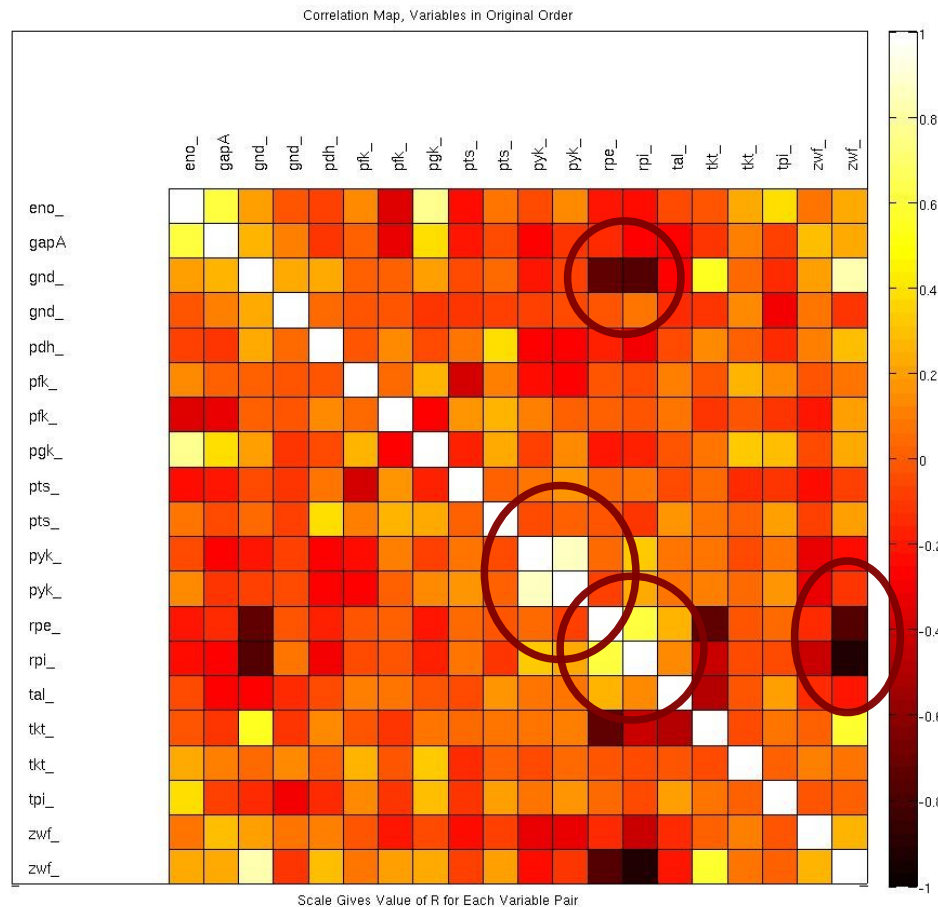
$$CC_p^x = \frac{p}{x(t)} \frac{\partial x}{\partial p}(t)$$



Several snapshots at different time points

Correlation among parameters

Correlation among parameters Based on Fischer Information Matrices



Strongly
Correlated
Parameters

Testing AD

Finite Difference Methods

$$\frac{\partial x}{\partial p_j}(t, p) = \frac{x(t, p + e_j \delta_j) - x(t, p)}{\delta_j} + O(\delta_j)$$

$$\delta_j = \epsilon p_j$$

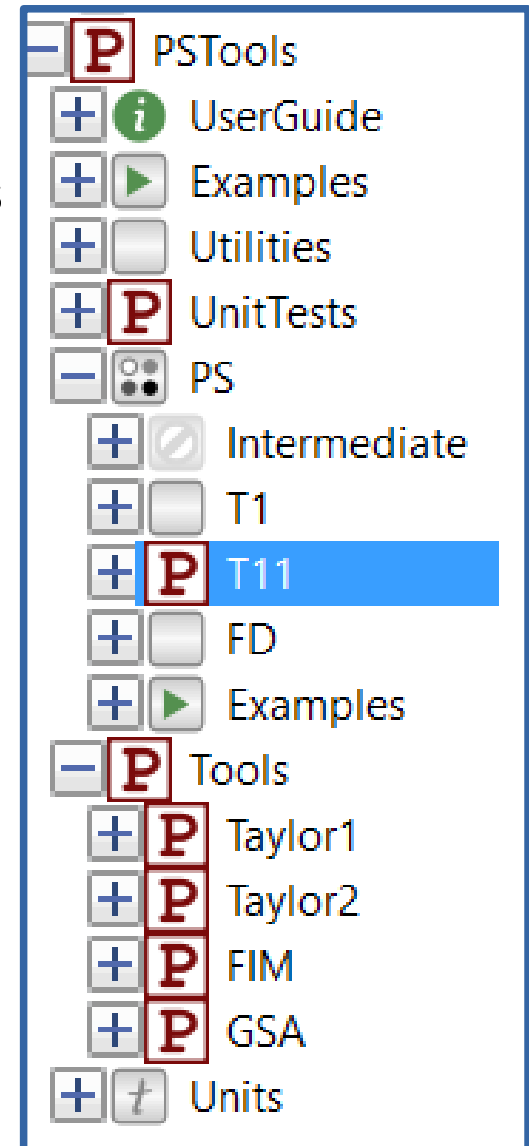
Adhoc Testing (2007-2014)

- 1) Evaluate DPS with FD
externally with Matlab
small models
- 2) Simulate AD models
- 3) Check if the curves behave similarly
Sometimes compute Relative errors
- 4) Investigate cause of errors, if any

PSTools Library

- Promotes the usage of PS at Modelica level
- Serves as utility package for arbitrary Modelica libraries
- PS Package
 - Extensive set of examples for analytical derivatives
 - Including hybrid systems
 - Generic Models for advanced FD
 - Second-order DPS
- Tools Package
 - Taylor series approximation
 - Parameter Sweeping studies
 - Control Coefficients
 - ...

Unit Testing is must



FD with PSTools

```
model ParM
```

```
  extends Utilities.Parameterized(
```

```
    NP = 2,
```

```
    _P = {0.4, 0.5},
```

```
    _PNAME = {"p1", "p2"},
```

```
    NX = 2,
```

```
    _X = {_M.x1, _M.x2},
```

```
    _XNAME = {"x1", "x2"};
```

```
);
```

```
protected
```

```
  MyModel _M(
```

```
    p1 = _P[1],
```

```
    p2 = _P[2]);
```

```
end ParM;
```

```
model FDParm
```

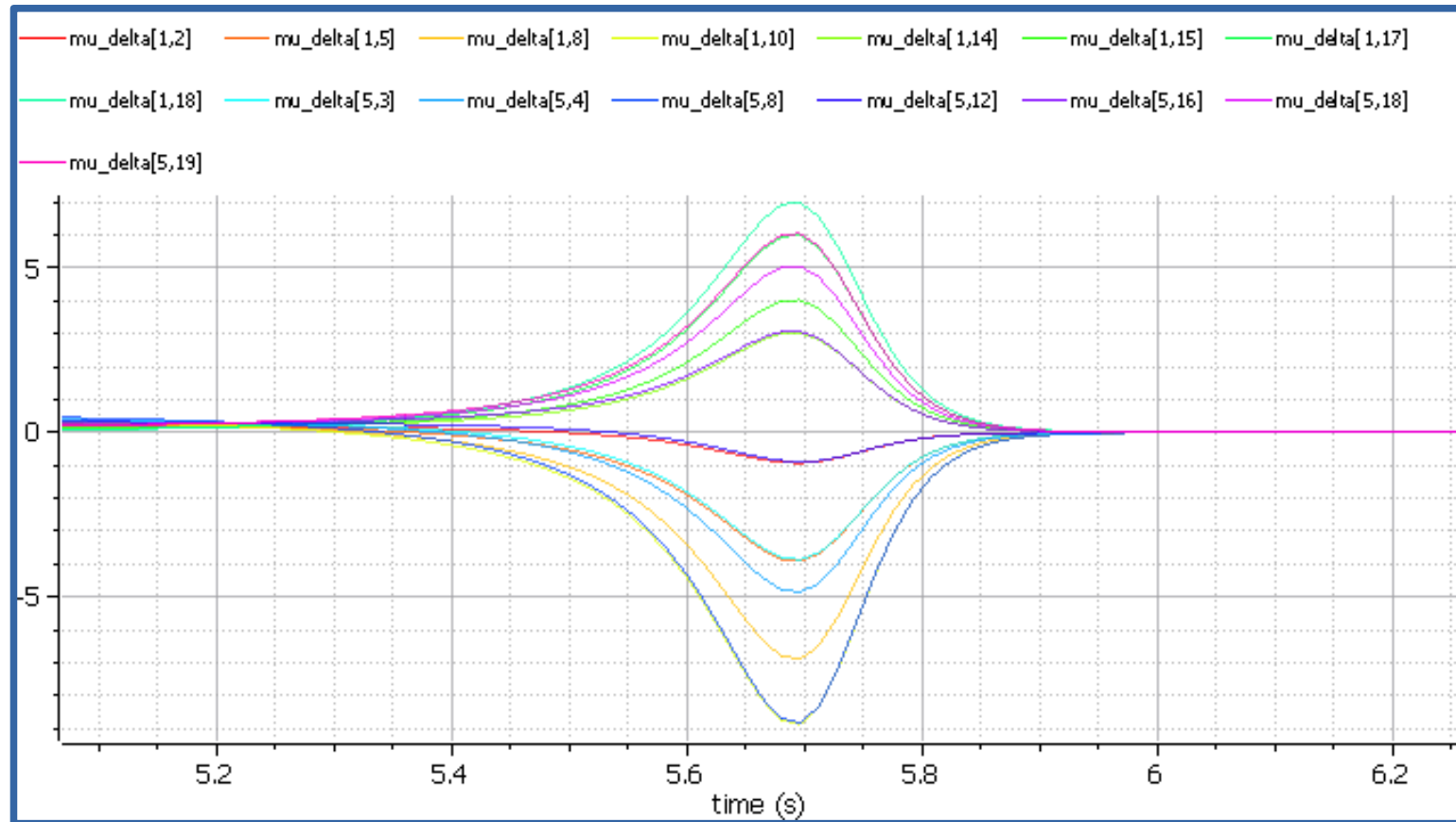
```
  PSTools.PS.FD.CD2
```

```
  PS(redeclare replaceable
```

```
    Model Parmodel = Parm);
```

```
end FDParm;
```

Parameter Sweeping Studies



Jan Peter Axelsson & Atiyah Elsheikh
Example of Sensitivity Analysis with the Bioprocess Library
Modprod 2019

Unit Testing Technologies for Modelica

Marco Kessler, Testing Tutorial
Dassault Systems, Modelica 2017 Prague ([Link](#))

- CSV Compare – ESI ITI
- BuildingPy – LBNL
- PySimulator – DLR
- test.openmodelica.org
- XogenyTest – Xogeny

- Model Management – Dymola
- Testing tool kits (many Companies)
- Testing Library -- Dymola

XogenyTest

- Pure Modelica
- Minimal
 - external scripting
 - log files
- Easy to use
- No dependencies

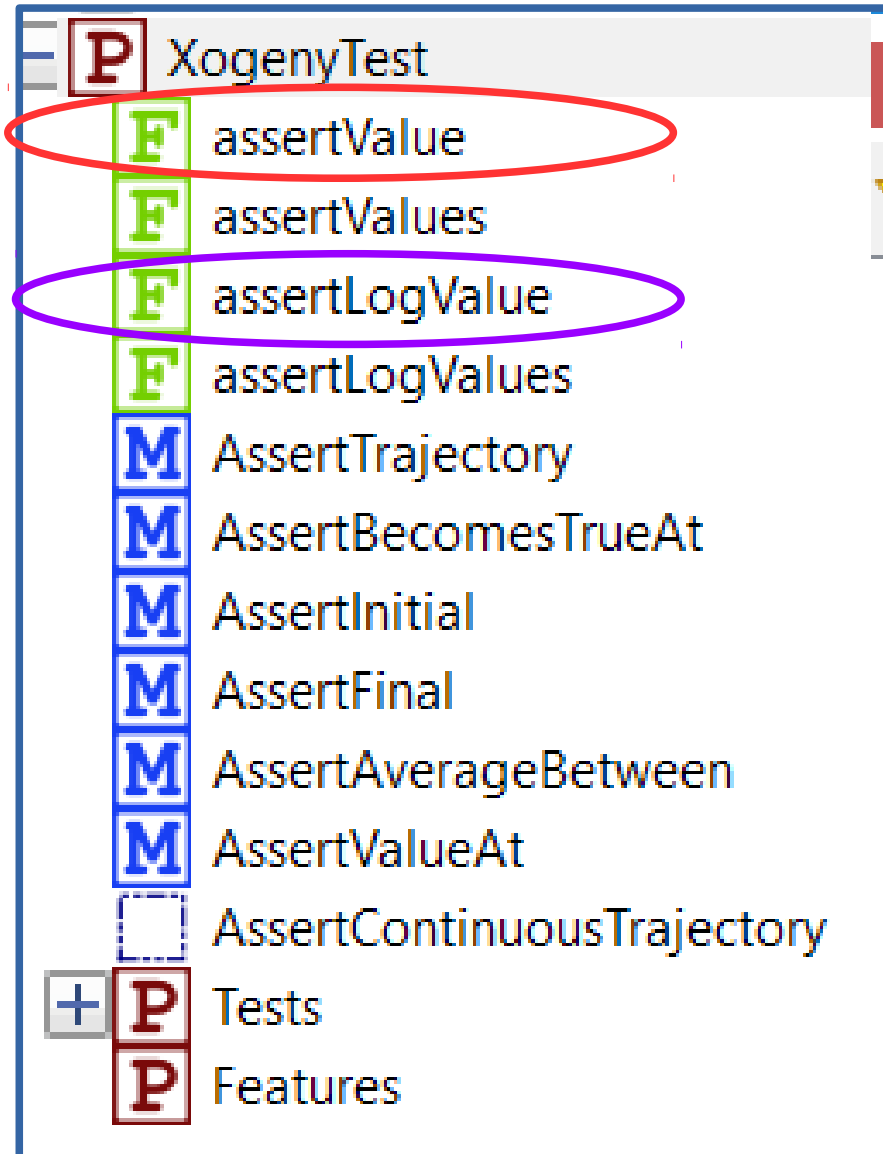
Not meant to be comprehensive but definitely an ideal getting start

Michael Tiller

<https://github.com/xogeny/XogenyTest>

Tiller, Michael M., and Burit Kittirungsi.
"UnitTesting: A Library for Modelica Unit Testing"
Modelica Conference Vienna, Austria, 2006

XogenyTest



Low level functions

asserting whether
an expected value x lies within
 ϵ -neighbourhood of a reference
value r (i.e. $N_\epsilon(r)$)

$$|x - r| < \epsilon$$

or

within the same order of
magnitude

$$\left| \log \frac{x}{r} \right| < \epsilon$$

XogenyTest

- P** XogenyTest
 - F** assertValue
 - F** assertValues
 - F** assertLogValue
 - F** assertLogValues
 - M** AssertTrajectory
 - M** AssertBecomesTrueAt
 - M** AssertInitial
 - M** AssertFinal
 - M** AssertAverageBetween
 - M** AssertValueAt
 - AssertContinuousTrajectory
- + P** Tests
- P** Features

Models

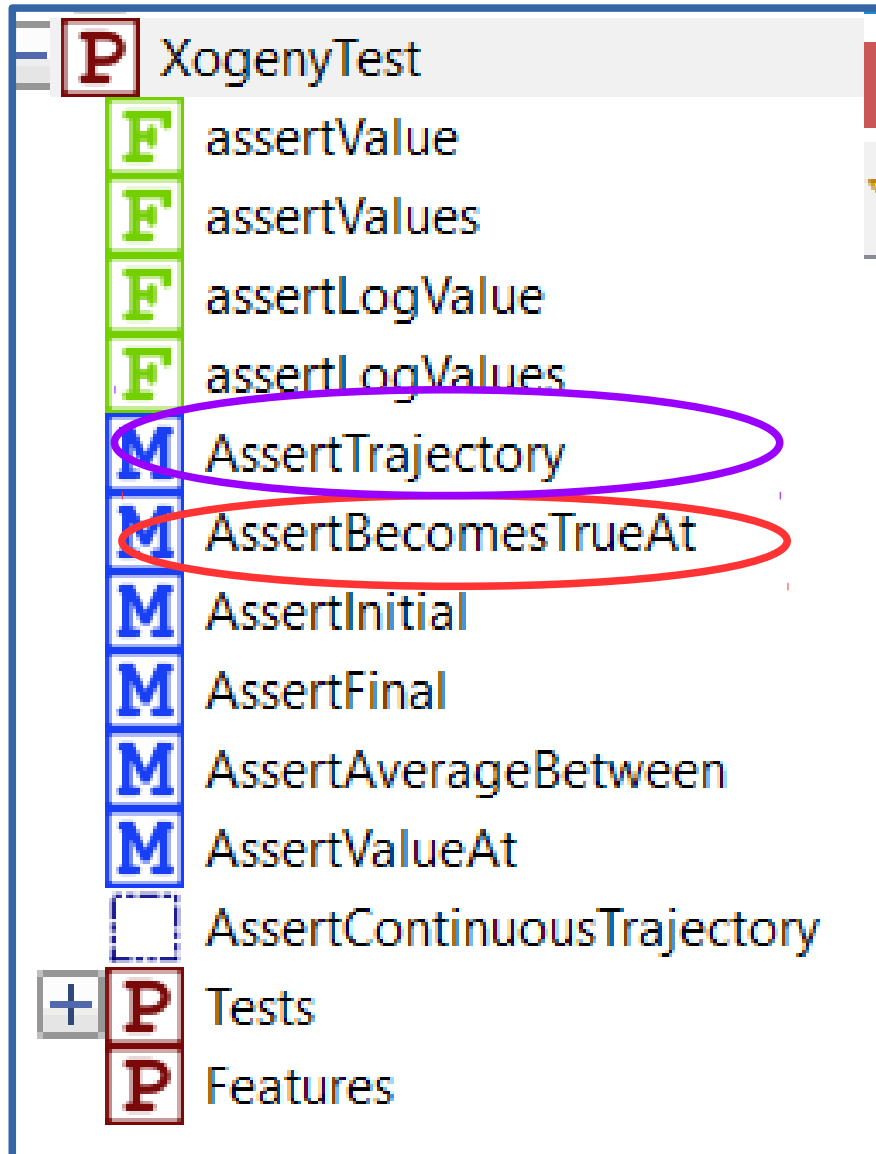
asserting an expected value x w.r.t. a reference value r at time $T \in \{t_0, t_f\}$

$$|x(T) - r| < \epsilon$$

or at time $t \in (t_0, t_f)$

$$|x(t) - r| < \epsilon$$

XogenyTest



Models

asserting an expected event $e(t)$ occurs at time T

$$e([T - \epsilon, T + \epsilon]) \rightarrow \text{True}$$

Asserting a trajectory $x(t)$ at discrete time points $\{t_1, t_2, \dots, t_N\}$ w.r.t. reference values $r(t_i)$

$$|x(t_i) - r(t_i)| < \epsilon$$

Examples (I/II)

```
model CheckSuccess
```

```
Real x = time^2;
```

```
AssertTrajectory check_x(  
  actual=x,  
  expected=[0,0; 1,1; 2,4; 3,9]);
```

```
Annotation(  
  TestCase(  
    action="simulate",  
    result="success"),  
  experiment(StopTime=4));
```

```
end CheckSuccess;
```

Examples (II/II)

```
model CheckSuccess
```

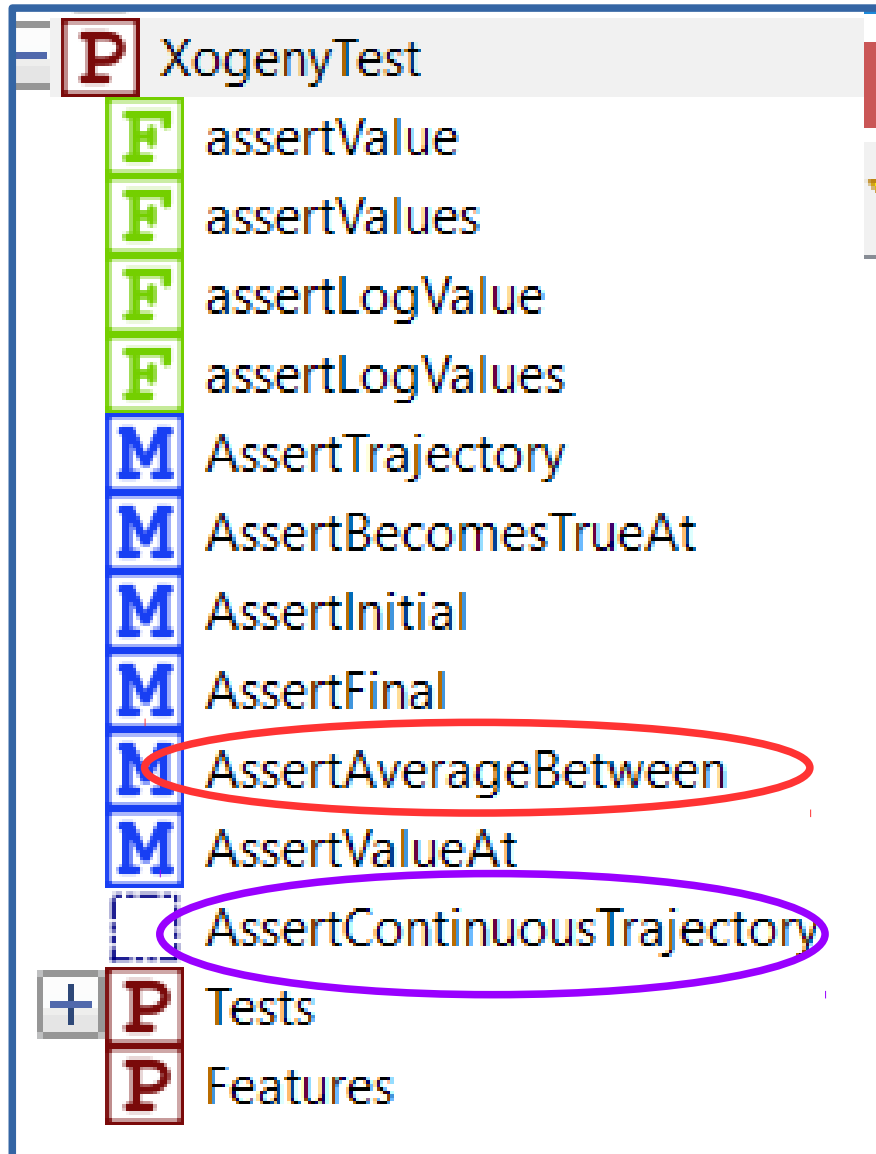
```
  Real x = time;
```

```
  AssertBecomesTrueAt  
    check_event(event=(x>2), at=2);
```

```
  Annotation(  
    TestCase(  
      action="simulate",  
      result="success"),  
    experiment(StopTime=4));
```

```
end CheckSuccess;
```

XogenyTest



Models

Assert that a signal $u(t)$ is of an average value a over $[t_0, t_f]$

$$\frac{1}{t_f - t_0} \left| \int_{t_0}^{t_f} u(t) dt \right| - a < \epsilon$$

Assert that difference between trajectories $x_1(t)$ and $x_2(t)$ is less than accu. error A_{err}

$$\int_{t_0}^{t_f} |x_1(t) - x_2(t)| dt < A_{err}$$

Testing AD vs. FD

```
model TestT1BioProcess
import PSTools.Utilities.unitVector;

Utilities.Validate dmu_dKs(
    AccErr=1E-1,
    name="test T1.Processions dmu / dKs");

PSTools.PS.T1.BioProcess.Bioprocess PSAD(
    NG=1,
    g_Ks=unitVector(1, NG));

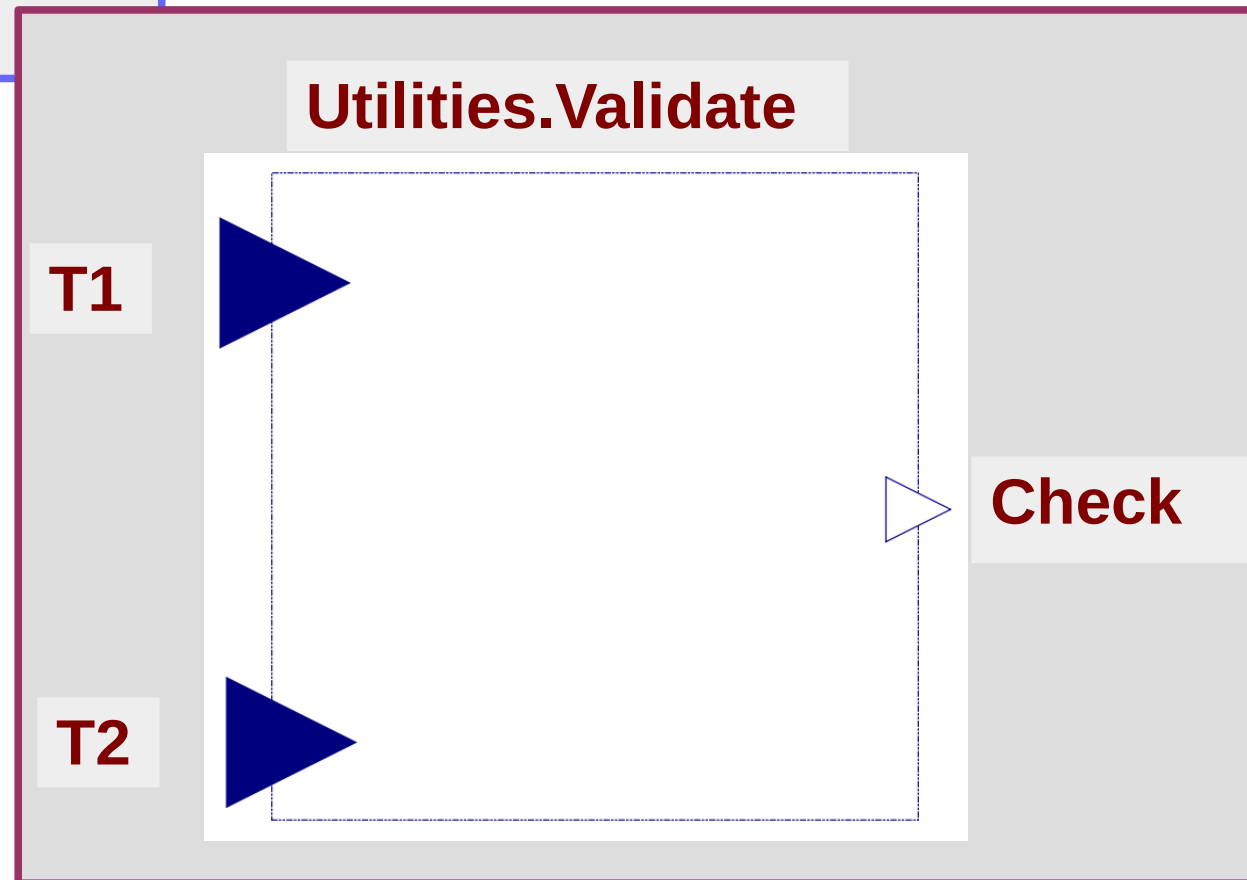
PSTools.PS.FD.BioProcess.BioProcess PSFD;
...
```

Visual unit testing

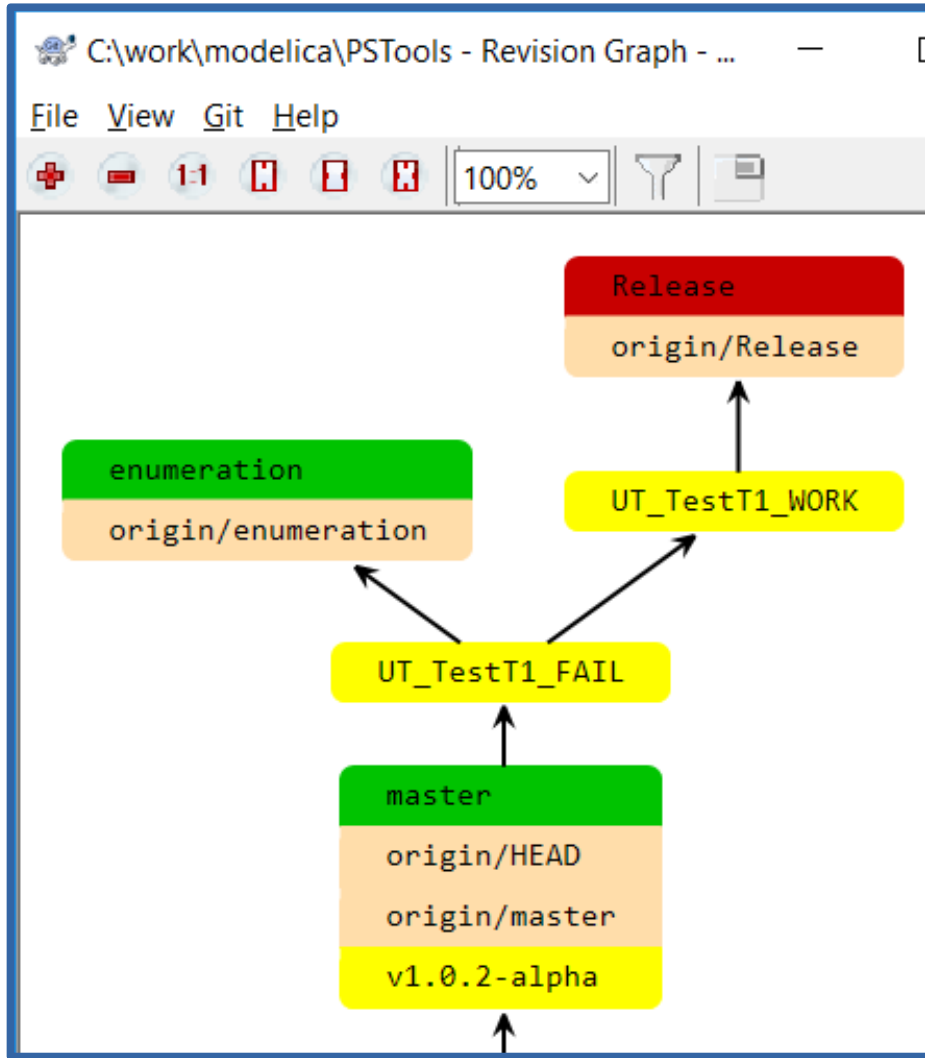
equation

```
dmu_dKs.T1 = PSAD.g_mu[1];  
dmu_dKs.T2 = PSFD.g_mu[1];
```

end TestT1BioProcess;

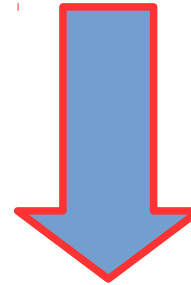


Case Study 1 PSTools Library



UT_TestT1_FAIL

```
// der(VS) = -qS * VX;  
g_VS[i] = -g_qS[i] * VX  
          - qS * g_VX[i];
```

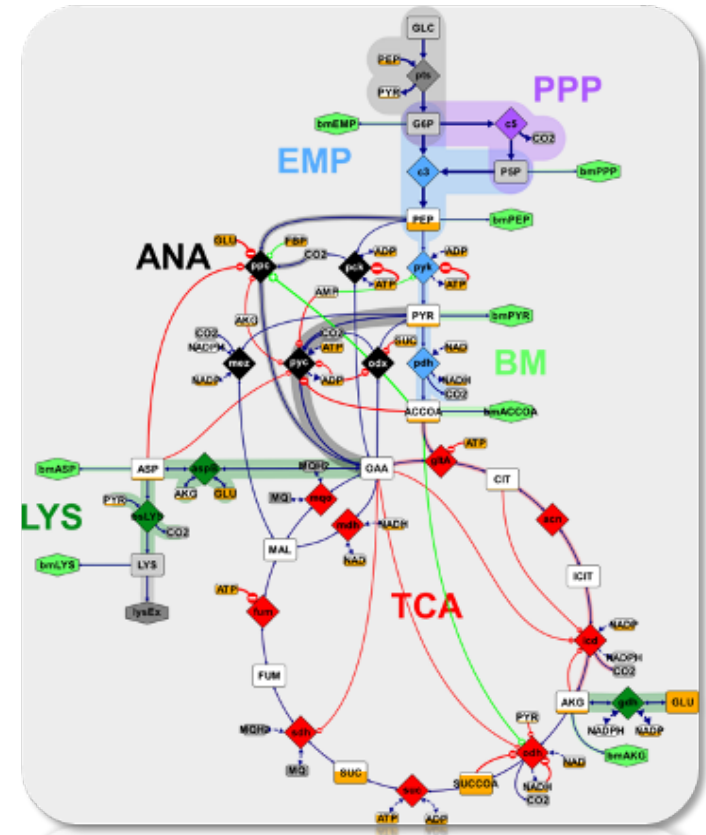
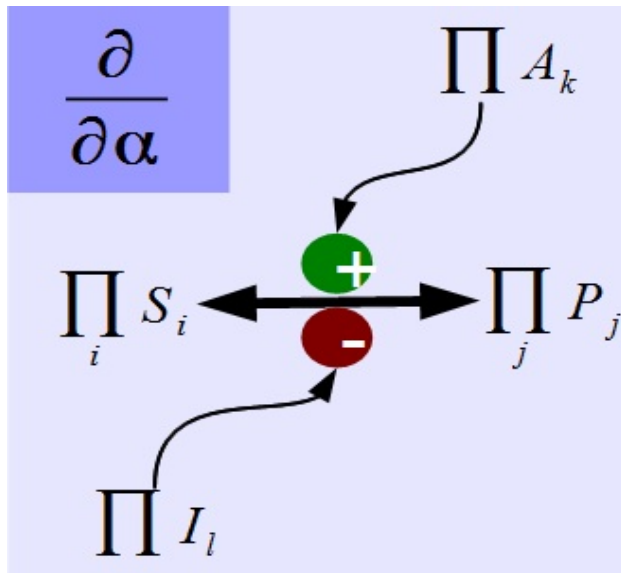


UT_TestT1_WORK

```
// der(VS) = -qS * VX;  
der(g_VS[i]) = -g_qS[i] * VX  
                - qS * g_VX[i];
```

GenKinetics

2018 R 1.0-alpha



$$v = \prod_a \frac{K_{A_a} + [A_a]}{K_{A_a}} \cdot \prod_b \frac{K_{I_b}}{K_{I_b} + [I_b]} \cdot \frac{V_{max}^{fwd} \prod_i \frac{[S_i]}{K_{mS_i}} - V_{max}^{bwd} \prod_j \frac{[P_j]}{K_{mP_j}}}{V_{max}^{fwd} \prod_i \left(1 + \frac{[S_i]}{K_{mS_i}}\right) + V_{max}^{bwd} \prod_j \left(1 + \frac{[P_j]}{K_{mP_j}}\right) - 1}$$

Formal unit testing procedure

- 1) a unit test model for each component C using
 - Example of a small model employing C
 - Equivalent model implemented only by equations
- 2) Execute all unit tests by *.mos & OMShell
 - after each significant modification
- 3) Investigate errors , if any

Most useful when the component checks but the UT does not translate or simulate

A library is shipped with its unit tests

All unit tests are within a package called UnitTests

Currently about 40 unit tests distributed GenKinetics

Case Study 2 ADGenKinetics Library

OMC 1.12.0

```
A = product({KA[i] / (KA[i] + mc_A[i].c) for i in 1:NA});
```

2012



OMC 1.13.0

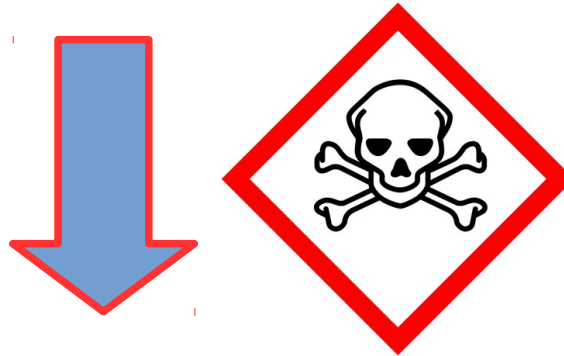
```
A = product({ (KA .+ mc_A.c) ./ KA for i in 1:NA});
```

2019

Case Study 3 GenKinetics / Biochem Libraries

Numerical Problem

```
der(c) = if  
  (c < tolerance) then 0 else r_net;
```



Solution

```
der(c) = if  
  (c < tolerance and r_net < 0) then 0 else r_net;
```

Advantages

- Simplicity
- Purely Modelica
- Suitability

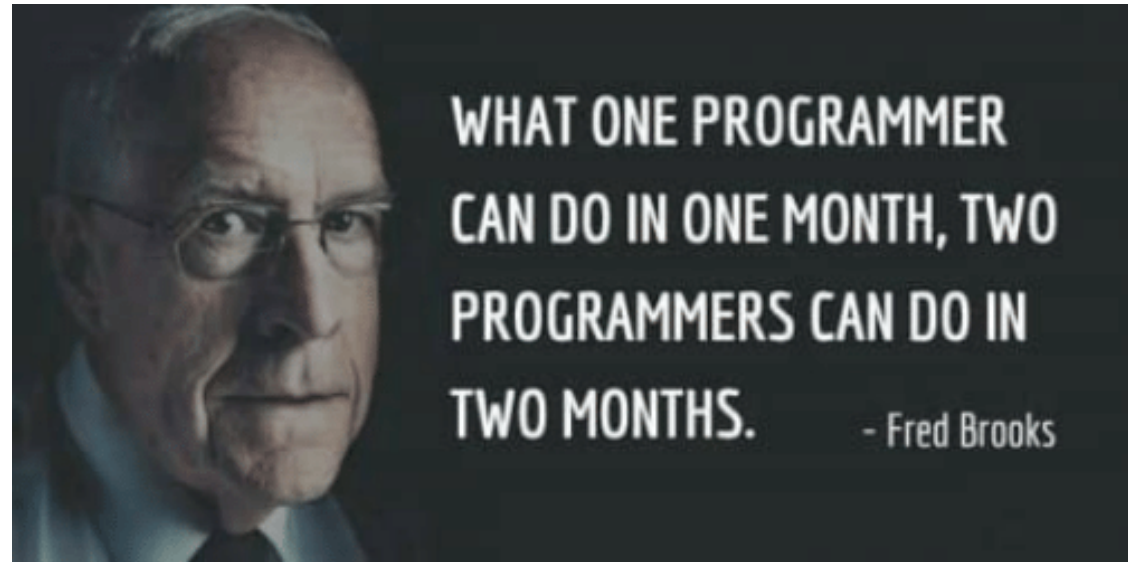
– individual developers or Modelica libraries

- GenKinetics <https://github.com/AtiyahElsheikh/GenKinetics>
- PSTools

– small-scale developments

– Minimum constraints get done

– Shipping Unit Tests with the product



Outlook

- a user experience report from GenKinetics
- Combine Relative Error w. Absolute Error

Asgar, Adeel, Andreas Pfeiffer, Arunkumar Palanisamy, Alachew Mengist, Martin Sjölund, Adrian Pop, and Peter Fritzson.

"Automatic Regression Testing of Simulation Models and Concept for Simulation of Connected FMUs in PySimulator."

In Proceedings of the 11th International Modelica Conference, Versailles. 2015.

- Continuous Integration Solution