

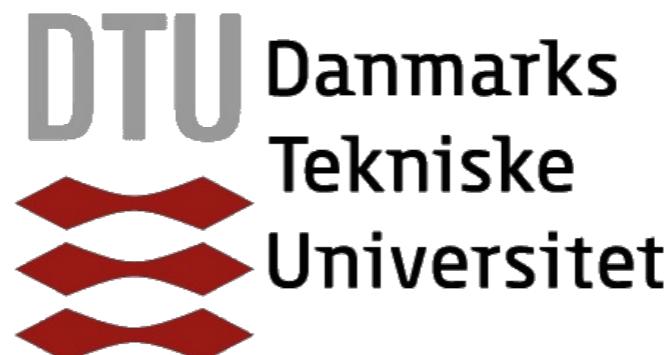
ERODE: EVALUATION AND REDUCTION OF DIFFERENTIAL EQUATIONS, CHEMICAL REACTION NETWORKS, BOOLEAN NETWORKS

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Department
of Excellence
2018 - 2022

EMbeDS
Economics and Management
in the era of Data Science



Based on joint work with

Luca Cardelli



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SCHOOL
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STUDIES
LUCCA

Max Tschaikowski



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But also

Tabea Waizmann, Isabel Perez Verona, Giuseppe Squillace, IMT Lucca

Stefano Tognazzi, University of Konstanz

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Giorgio Bacci, Giovanni Bacci, Kim G. Larsen, Aalborg University

Nicolas Gast, INRIA

Luca Bortolussi, University of Trieste

ERODE: OVERVIEW OF THE TOOL'S FUNCTIONALITIES

The screenshot displays the ERODE software interface, which includes:

- Project Explorer:** Shows a file tree with "Examples" containing "ExampleODE.ode" and "ExampleRN.ode".
- Code Editors:** Two editors show the model definitions:
 - ExampleODE.ode:** A chemical reaction system with parameters r1=1.0, r2=2.0, and species Au, Ap, B, AuB, ApB. It includes ODEs for the species concentrations.
 - ExampleRN.ode:** A reduced version of the model with reactions Au → Ap, Ap → Au, Au + B → AuB, and AuB → Au + B.
- Outline View:** Provides a hierarchical overview of the code elements.
- Console:** Displays the command-line output of the simulation:

```
ERODE -ExampleRN-[15/05/2016 18-57-46-218]
*****
***** ERODE -ExampleRN-[15/05/2016 18-57-46-218] *****
*****
Reading ExampleRN...
Parameters: 2
Species: 5
Reactions: 6.

Solving ODEs of ExampleRN... completed in 0.006 (s).
```
- Simulation Results:** A graph titled "simulateODE(tEnd=1.0)" showing the concentration of five species over time (Time from -0.01 to 1.01). The species are: Au (blue), Ap (red), B (green), AuB (black), and ApB (purple). The concentrations stabilize after approximately 0.5 units of time.
- Logo:** A circular logo for "ERODE" with the text "Evaluation and Reduction of Ordinary Differential Equations" around the border.

[TACAS17]

ERODE: OVERVIEW OF THE TOOL'S FUNCTIONALITIES

The screenshot shows the ERODE software interface with several open model files:

- AM_RN.ode**:

```
begin model AM_RN
begin parameters
p1=1 p2=2 p3=3 p4=4
end parameters
begin init
x0 = 100 x1 x2 = 100
end init
begin reactions
x0 + x2 -> x2 + x1 , p1
x0 + x2 -> x0 + x1 , p2
x0 + x1 -> x0 + x0 , p3
x2 + x1 -> x2 + x2 , p4
end reactions
//Analysis
simulateODE(tEnd=0.05)
simulateCTMC(tEnd=1,
repeats=100)
//Reduce
reduceBE()    reduceFE()
//Export
write(fileOut="AM_ODE.ode",
format=ODE)
exportBNG(
fileOut="AM_RN.net")
exportSBML(
fileOut="AM_RN.sbml")
exportStochKit(
fileOut="AM_RN.xml")
exportLNA(fileOut="LNA.ode")
generateCME(
fileOut="CME.ode")
end model
```
- AM_ODE.ode**:

```
begin model AM_ODE
begin parameters
p1 = 1 p2 = 2 p3 = 3 p4 = 4
end parameters
begin init
x0 = 100 x1 x2 = 100
end init
begin ODE
d(x0) = p3*x0*x1 - p1*x0*x2
d(x1) = (p1+p2)*x0*x2 - p3*x0*x1
- p4*x1*x2
d(x2) = p4*x1*x2 - p2*x0*x2
end ODE
//Analysis
simulateODE(tEnd=0.05,
library=APACHE)
simulateODE(tEnd=1,
library=SUNDIALS)
//Reduction
reduceBDE()
//Export
reduceFDE(reducedFile=
"AM_ODE_FDE.ode")
//Export
exportMatlab(fileOut="AM_ODE.m",
tEnd=1)
end model
```
- AM_DAE.ode**:

```
begin model AM_DAE
begin parameters
p1 = 1 p2 = 2 p3 = 3 p4 = 4
end parameters
begin init
x0 = 100 x1
end init
begin alginit
x2 = 100
end alginit
begin ODE
d(x0) = p3*x0*x1 - p1*x0*x2
d(x1) = (p1+p2)*x0*x2 - p3*x0*x1
- p4*x1*x2
end ODE
begin algebraic
x2 = 200 - x0 - x1
end algebraic
simulateDAE(tEnd=1)
exportModelica(
fileOut="AM_DAE.mo")
reduceBDE()
end model
```
- *BN.ode**:

```
begin Boolean Network BN
begin init
x1=true x2=false x3=true
end init
begin update functions
x1 = ! x3 or x1
x2 = x1 or x2 or ! x3
x3 = x2 and ! x3
end update functions
reduceBBE(reducedFile="BN_BBE.ode")
end Boolean Network
```
- *BN_BBE.ode**:

```
//Generated from BN via BBE
//Size of initial partition: 1
//Original number of species: 3
//Reduced number of species: 2
begin Boolean network BN_BBE
begin init
x1 = true
x3 = true
end init
begin update functions
x1 = ((!x3) | x1)
x3 = (x1&(!x3))
end update functions
//Comments associated to the species
//x1:
//Representative of block
// x1,x2
//x3:
//Singleton block
end Boolean network
```
- importers.ode**:

```
begin model mrmc
importMRMC(fileIn="ctmc.tra",
labellingFile="ctmc.lab")
end model
```
- importers2.ode**:

```
begin model affine
importAffineSystem(
fileIn="A.csv", BFile="b.csv",
ICFile="IC.csv")
end model
```
- importers3.ode**:

```
begin model bng
importBNG(fileIn="bng.net")
end model
```

ERODE: OVERVIEW OF THE TOOLS FUNCTIONALITIES

runtime-ERODE.product(4) - PER/AM_ODE.ode - ERODE

The screenshot shows the ERODE software interface with multiple code editors and a plot window.

Code Editors:

- AM_RN.ode:**

```
begin model AM_RN
begin parameters
p1=1 p2=2 p3=3 p4=4
end parameters
begin init
x0 = 100 x1 x2 = 100
end init
begin reactions
x0 + x2 -> x2 + x1 , p1
x0 + x2 -> x0 + x1 , p2
x0 + x1 -> x0 + x0 , p3
x2 + x1 -> x2 + x2 , p4
end reactions
//Analysis
simulateODE(tEnd=0.05)
simulateCIMC(tEnd=1,
repeats=100)
//Reduce
reduceBE()
reduceFE()
//Export
write(fileOut="AM_ODE.ode",
format=ODE)
exportBNG(
fileOut="AM_RN.net")
exportSBML(
fileOut="AM_RN.sbml")
exportStochKit(
fileOut="AM_RN.xml")
exportLNA(fileOut="LNA.ode")
generateCME(
fileOut="CME.ode")
end model
```
- AM_ODE.ode:**

```
begin model AM_ODE
begin parameters
p1 = 1 p2 = 2 p3 = 3 p4 = 4
end parameters
begin init
x0 = 100 x1 x2 = 100
end init
begin ODE
d(x0) = p3*x0*x1 - p1*x0*x2
d(x1) = (p1+p2)*x0*x2 - p3*x0*x1
- p4*x1*x2
d(x2) = p4*x1*x2 - p2*x0*x2
end ODE
//Analysis
simulateODE(tEnd=0.05,
library=APACHE)
simulateODE(tEnd=1,
library=SUNDIALS)
//Reduction
reduceBDEC()
reduceFDEC(reducedFile=
"AM_ODE_FDE.ode")
//Export
exportMatlab(fileOut="AM_ODE.m",
tEnd=1)
end model
```
- importers2.ode:**

```
begin model affine
importAffineSystem(
fileIn="A.csv", BFile="b.csv",
ICFile="IC.csv")
end model
```

Plot Window:

AM_ODE - ODE solutions

The plot shows three variables over time (0 to 0.05). The x-axis is labeled "Time" and the y-axis is labeled "Variable concentrations".

Time	x0 (Blue)	x1 (Red)	x2 (Green)
0.000	100	0	100
0.005	110	35	65
0.010	130	30	40
0.015	150	20	30
0.020	170	10	20
0.030	190	5	15
0.040	195	2	12
0.050	198	0	10

Legend: x0 (Blue), x1 (Red), x2 (Green)

ERODE: OVERVIEW OF THE TOOL'S FUNCTIONALITIES

The screenshot shows the ERODE software interface with two code editors. The top editor is titled "runtime-ERODE.product(4) - PER/AM_ODE.ode - ERODE" and contains the following code:

```
begin Boolean Network BN
begin init
    x1=true x2=false x3=true
end init
begin update functions
    x1 = ! x3 or x1
    x2 = x1 or x2 or ! x3
    x3 = x2 and ! x3
end update functions
reduceBBE(reducedFile="BN_BBE.ode")
end Boolean Network
```

The bottom editor is titled "*BN_BBE.ode" and contains the generated code:

```
//Generated from BN via BBE
//Size of initial partition: 1
//Original number of species: 3
//Reduced number of species: 2
begin Boolean network BN_BBE
begin init
    x1 = true
    x3 = true
end init
begin update functions
    x1 = ((!x3) | x1)
    x3 = (x1&(!x3))
end update functions
//Comments associated to the species
//x1:
//Representative of block
//    x1,x2
//x3:
//Singleton block
end Boolean network
```

ERODE: OVERVIEW OF THE TOOLS FUNCTIONALITIES

runtime-ERODE.product(5) - tool_demo/SIRRNHull_union.ode - ERODE

The screenshot shows the ERODE graphical user interface with three code editors and a plot window.

- LNA.ode:**

```

begin model LNA
begin parameters
p1 = 1.0 p2 = 2.0 p3 = 3.0 p4 = 4.0
end parameters
begin init
x0 = 100.0 x1 x2 = 100.0
C_x0_x0 C_x0_x1 C_x0_x2
C_x1_x0 C_x1_x1 C_x1_x2
C_x2_x0 C_x2_x1 C_x2_x2
end init
begin ODE
d(x0) = -p1*x0*x2 + p3*x0*x1
d(x1) = p1*x0*x2 + p2*x0*x2 + -p3*x0*x1 + -p4*x1*x2
d(x2) = -p2*x0*x2 + p4*x1*x2
d(C_x0_x0) = (-p1*x2 + p3*x1) * C_x0_x0 + p3*x0 * C_x0_x1
d(C_x0_x1) = (-p1*x2 + p3*x1) * C_x0_x1 + p3*x0 * C_x0_x2
d(C_x0_x2) = (-p1*x2 + p3*x1) * C_x0_x2 + p3*x0 * C_x1_x0
d(C_x1_x0) = (p1*x2 + p2*x2 - p3*x1) * C_x0_x0 + (-p1*x1 + p3*x2) * C_x0_x1
d(C_x1_x1) = (p1*x2 + p2*x2 - p3*x1) * C_x0_x1 + (-p1*x1 + p3*x2) * C_x0_x2
d(C_x1_x2) = (p1*x2 + p2*x2 - p3*x1) * C_x0_x2 + (-p1*x1 + p3*x2) * C_x1_x0
d(C_x2_x0) = -p2*x2 * C_x0_x0 + p4*x2 * C_x1_x0 + (-p2*x1 * C_x0_x1 + p4*x1 * C_x1_x0)
end ODE

```
- SIR2.ode:**

```

begin model SIR2
begin parameters
b11 = 1.04 b12 = 0.96
b21 = 1.05 b22 = 0.95
g1 = 0.1 g2 = 0.1
end parameters
begin init
s1 = 0.48 s2 = 0.51
i1 = 0.52 i2 = 0.49
r1 r2
end init
begin ODE
d(s1) = -b11*s1*i1 + -b12*s1*i2
d(s2) = -b21*s2*i1 + -b22*s2*i2
d(i1) = b11*s1*i1 + b12*s1*i2 + -g1*i1
d(i2) = b21*s2*i1 + b22*s2*i2 + -g2*i2
d(r1) = g1*i1
d(r2) = g2*i2
end ODE

```
- SIR2Hull.ode:**

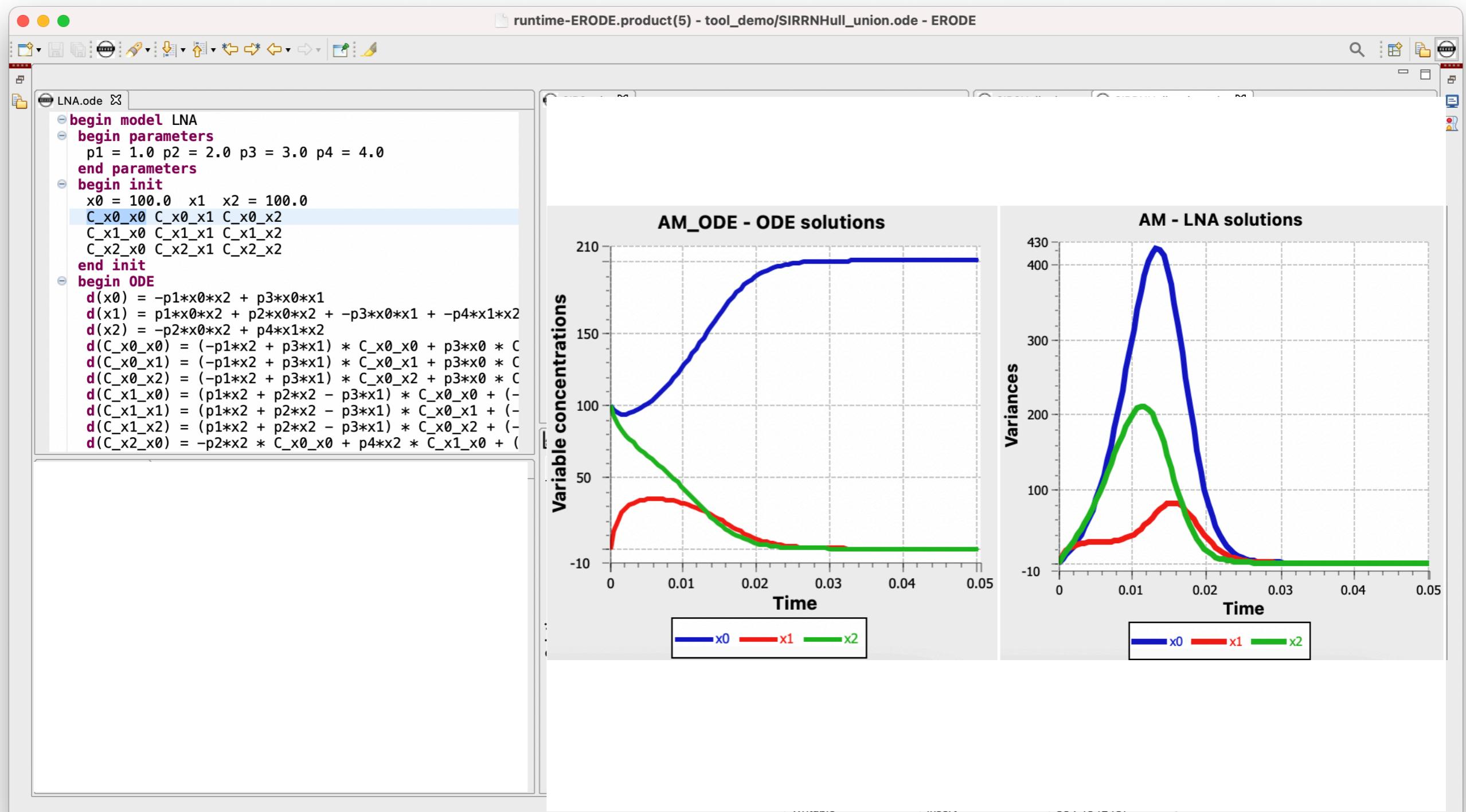
```

begin model SIRRNHull_union
begin parameters
b11=1.04 b12=0.96 b21=1.05 b22=0.95
g1=0.1 g2=0.1
end parameters
begin init
us1 = 0.48 os1 = 0.48
us2 = 0.51 os2 = 0.51
ui1 = 0.52 oi1 = 0.52
ui2 = 0.49 oi2 = 0.49
ur1 or1
ur2 or2
s1 = 0.48 s2 = 0.51
i1 = 0.52 i2 = 0.49
r1 r2
end init
begin ODE
d(us1)= -b21*us1*oi1 - b12*us1*oi2
d(os1)= -b11*os1*ui1 - b22*os1*ui2
d(us2)= -b21*us2*oi1 - b12*us2*oi2
d(os2)= -b11*os2*ui1 - b22*os2*ui2
d(ui1)= b11*us1*ui1 + b22*us1*ui2 - g1*ui1
d(oi1)= b21*os1*oi1 + b12*os1*oi2 - g1*oi1
d(ui2)= b11*us2*ui1 + b22*us2*ui2 - g1*ui2
d(oi2)= b21*os2*oi1 + b12*os2*oi2 - g1*oi2
d(ur1)=
d(or1)=
d(ur2)=
d(or2)=
d(s1) = -b11*s1*i1 - b12*s1*i2
d(s2) = -b21*s2*i1 - b22*s2*i2
d(i1) = b11*s1*i1 + b12*s1*i2 - g1*i1
d(i2) = b21*s2*i1 + b22*s2*i2 - g2*i2
d(r1) =
d(r2) =
end ODE

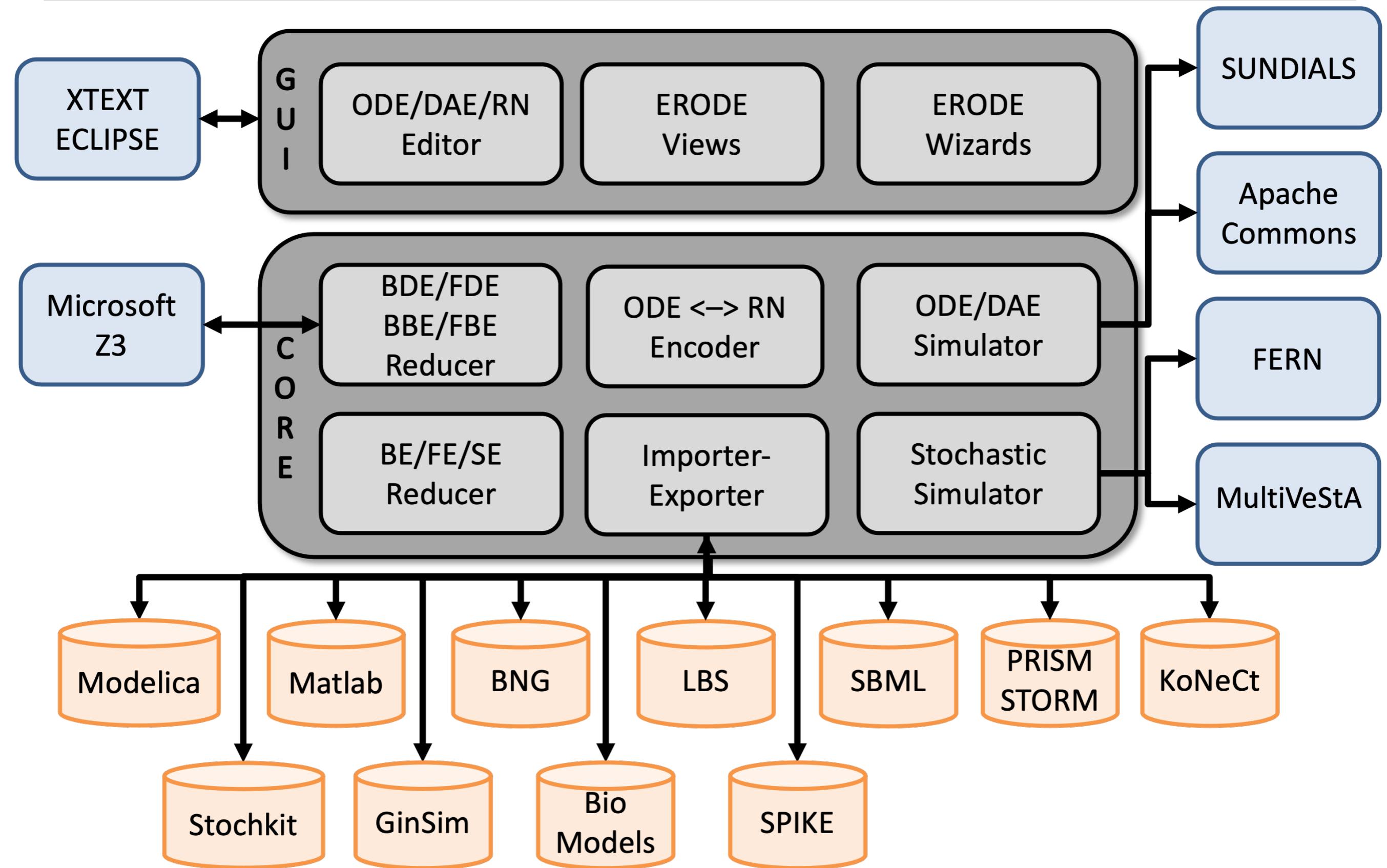
```

Plot Window: The plot window displays the "Solution of the hull and of the original model" over Time (0 to 5.05). It shows three curves: *i1* (solid blue), *o_i1* (dashed red), and *u_i1* (dash-dot green).

ERODE: OVERVIEW OF THE TOOLS FUNCTIONALITIES



ERODE: ARCHITECTURE AND SOME OF THE SUPPORTED I/O FORMATS



SOME REFERENCES

TECHNICAL PRESENTATIONS

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FUNDING & ONGOING AND FUTURE WORK

- ▶ GRANTS with most support for the tool:
 - ▶ FEMPA, DFG, Germany
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 - ▶ COCO, FWF, Austria
 - ▶ REDUCTO, DFF RP1, Denmark
 - ▶ SEDUCE, PRIN, Italy
 - ▶ Danish PDJ foundation
- ▶ Ongoing and future work
 - ▶ Control-related reductions/analysis of networks (COCO)
 - ▶ Rethink the framework for Boolean networks (REDUCTO)
 - ▶ Alternative approaches to partition refinement [*LICS'21*]
 - ▶ All our algorithms so far are for exact reductions
 - ▶ Approximate reductions in terms of exact ones on perturbed models [*QEST'18*]
 - ▶ What if the model has intrinsic uncertainty? [*QEST'21*]
 - ▶



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www.erode.eu