

# On the Package ReactionKinetics.wl

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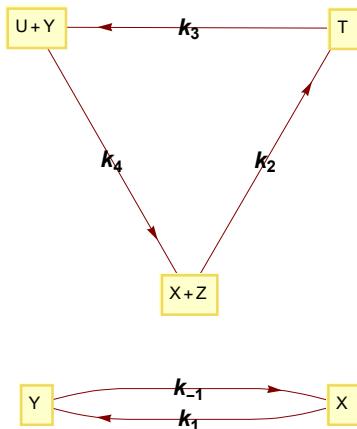
## Preliminaries

```
In[1]:= SetDirectory[NotebookDirectory[]];  
  
In[2]:= SetOptions[#, AxesStyle -> Arrowheads[Automatic]] & /@  
    {ContourPlot, DateListPlot, Plot, ListLinePlot, ListPlot, ListLogPlot,  
     LogLinearPlot, LogPlot, ParametricPlot, Plot3D, RegionPlot};  
SetOptions[GraphPlot, DirectedEdges -> True, VertexLabeling -> All];  
  
In[4]:= LaunchKernels[];  
  
In[5]:= Needs["ReactionKinetics`"]  
+  
ReactionKinetics Version 1.0 [March 25, 2018] using  
Mathematica Version 11.3.0 for Microsoft Windows (64-bit) (March 7,  
2018) (Version 11.3, Release 0) loaded 12 April 2018 at 11:05 TimeZone  
GNU General Public License (GPLv3) Terms Apply.  
Please report any issues, comments, complaint related to ReactionKinetics at  
jtoth@math.bme.hu, nagyal@math.bme.hu or dpapp@iems.northwestern.edu  
  
In[6]:= ClearAll[format, figureexp];  
format[x_, options___] :=  
    Thread[x -> (Style[#, SingleLetterItalics -> False, options] & /@ x)];  
figureexp[filename_, picturename_] := Export[  
    "d:\\__Springer\\SpringerAktualis\\Figures\\\" <> filename <> ".png", picturename]
```

# What is formal reaction kinetics?

## Feinberg-Horn III

```
ShowFHJGraph[FHIII = {"X" <=> "Y", "X" + "Z" -> "T" -> "Y" + "U" -> "X" + "Z"}, {k1, k-1, k2, k3, k4}, EdgeLabeled -> True, ImageSize -> 200] /. {kw -> Style[kw, Bold, 12]}
```

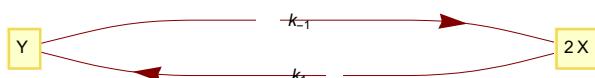


```
RightHandSide[{FHIII}, {k1, k-1, k2, k3, k4}, {x, y, z, t, u}] // Column
y k-1 - x k1 - x z k2 + u y k4
- y k-1 + x k1 + t k3 - u y k4
- x z k2 + u y k4
x z k2 - t k3
t k3 - u y k4
```

How to say anything about stationary points and dynamic behavior? Let us take a simpler example.

## Dimerization

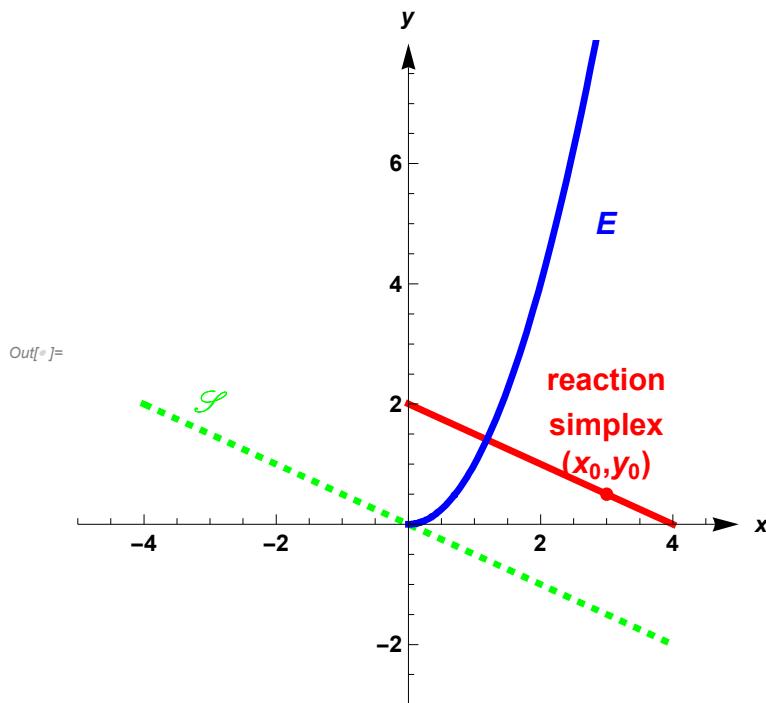
```
ShowFHJGraph[dimer = {2 "X" <=> "Y"}, {k1, k-1}, EdgeLabeled -> True]
```



```
Column@First@DeterministicModel[dimer, {k1, k-1}, {x, y}]
```

$$\begin{aligned} x'[t] &= -2 k_1 x[t]^2 + 2 k_{-1} y[t] \\ y'[t] &= k_1 x[t]^2 - k_{-1} y[t] \end{aligned}$$

```
In[=]:= statpoints = ParametricPlot[Evaluate[
  Flatten[{x, y} /. Rest[StationaryPoints[{2 "X" <=> "Y"}, {1, 1}, {u, v}, {x, y}]]] /.
  (Last[First[GammaLeftNullSpace[{2 "X" <=> "Y"}, {u, v}, {x, y}]]] /. Plus -> Rule)],
{v, 0, 10}, PlotStyle -> Directive[{Blue, Thickness[0.01]}],
Prolog -> {Directive[{Green, Dashed, Thickness[0.01]}],
  Line[{{{-4, 2}, {4, -2}}], Directive[{Red, Dashing[{0}], Thickness[0.01]}]
  , Line[{{0, 2}, {4, 0}}], PointSize[0.02], Point[{3, 1/2}], Green,
  Text[Style["S", FontFamily -> "Kunstler Script", Bold, 16], {-3, 2}],
  Red, Text[Style["reaction\`nsimplex", Bold, 16], {3, 2}],
  Text[Style["(x0,y0)", Bold, 16], {3, 1}], Blue,
  Text[Style["E", Italic, Bold, 16], {3, 5}]},
AspectRatio -> 1, PlotRange -> {{-5, 5}, {-3, 8}}},
AxesLabel -> (Style[#, Bold, Italic, 12] & /@ {"x", "y"}),
TicksStyle -> Directive[Black, Bold, 12]]]
```



Why does it behave so well?

```
ReactionsData[{2 "X" <=> "Y"}]["deficiency"]
```

$$\delta = N - L - S = 2 - 1 - 1 = 0$$

```
WeaklyReversibleQ[{2 "X" <=> "Y"}]
```

```
True
```

Suppose that the deficiency ( $N - L - S$ ) of a reaction is zero. Then:

1. No nontrivial periodic solutions of the induced kinetic differential equation can exist.
2. Furthermore,
  - a. If the reaction is not weakly reversible then it cannot have a positive stationary point, no matter what form the kinetic has.
  - b. If the reaction is weakly reversible and the kinetic is of the mass action form then with any choice of positive rate constants (i.e. with any mechanism built on the given reaction):

- i. There exists in each positive stoichiometric compatibility class exactly one positive stationary concentration.
  - ii. Each of the positive stationary concentrations is **relatively** asymptotically stable relative to the stoichiometric compatibility class in which it resides.
- 

## Returning to FH III

```
In[8]:= FHIII = {"X" <=> "Y", "X" + "Z" -> "T" -> "U" + "Y" -> "X" + "Z"};
```

```
WeaklyReversibleQ[{FHIII}]
```

```
True
```

```
ReactionsData[{FHIII}] ["deficiency"]
```

```
 $\delta = N - L - S = 5 - 2 - 3 = 0$ 
```

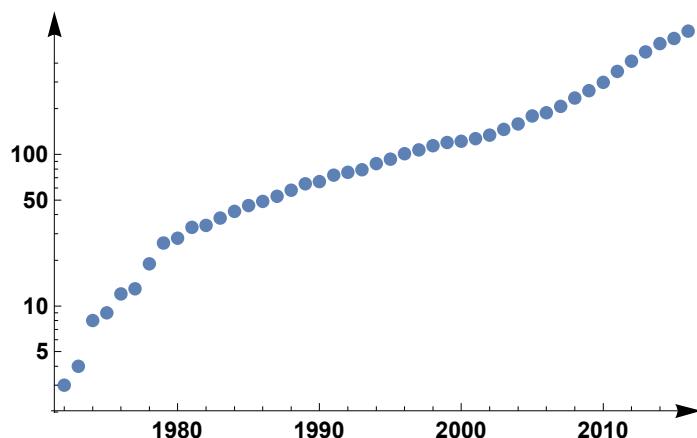
## Citations to the paper by Horn and Jackson

Data collected manually from scholar.

```
cit = {3, 4, 8, 9, 12, 13, 19, 26, 28, 33, 34, 38, 42, 46, 49,
      53, 58, 64, 66, 73, 76, 79, 87, 93, 101, 107, 114, 120, 122, 127, 134,
      146, 159, 179, 188, 207, 235, 263, 298, 353, 412, 475, 537, 581, 649};
```

The logarithm of the number of citations of Horn and Jackson (1972) up to a certain year. Note the three phases: Up to 1980, and after 2002 and in between.

```
ListLogPlot[Transpose[{Range[1972, 2016], cit}],
 PlotStyle -> Directive[PointSize[Large]], TicksStyle -> Directive[Black, Bold, 12]]
```



## Our early history

Stochastic simulation since 1972 (Érdi, P.; Hárs, V., Sipos, T.; Tóth, J.)

Mathematica codes for reaction kinetics since 2002 (Nagy A. L.; Nagy, I.; Papp, D.; Rospars, J.-P.; Tóth, J.)

and students :) such as Bányász, K.; Berencsi K.; Eszes, D.; Halschläger A.; Kovács, B.; Ladics, T.; Sipos Szabó E.; Szabó A.; Takács B.; Varga N.; Várdainé Kollár J.; Zsámboki B.; Vizi Zsolt (Szeged) )  
Numerics and beyond (or before): What can be done with the computer?

---

## Goals

Calculations of conditions of known theorems and of our results to help the chemist in case if they are not feasible manually.

Solving induced kinetic differential equations, simulating the stochastic model, estimating parameters.

## Physical circumstances

```
In[°]:= lvgenuine = ReactionsData[{"Lotka-Volterra"}];
In[°]:= lvgenuine["reactionsteps"]
Out[°]= {A + X → 2 X, X + Y → 2 Y, Y → B}

In[°]:= lv = ReactionsData[{"Lotka-Volterra"}, {"A", "B"}];
In[°]:= lv["reactionsteps"]
Out[°]= {A + X → 2 X, X + Y → 2 Y, Y → B}

In[°]:= lv[{"fhjgraphedges}]
Out[°]= {X → 2 X, X + Y → 2 Y, Y → 0}

lv["species"]
{X, Y}

lv["externalspecies"]
{A, B}
```

Constant pressure and volume; temperature may change. In- and outflow is represented using the empty complex.

Usual assumptions of the chemist (**mass conservation, low orders of reactions, reversibility, detailed balancing**) are not assumed from the beginning, they may be used as restrictions, their appearance can be checked using necessary and sufficient conditions, etc.

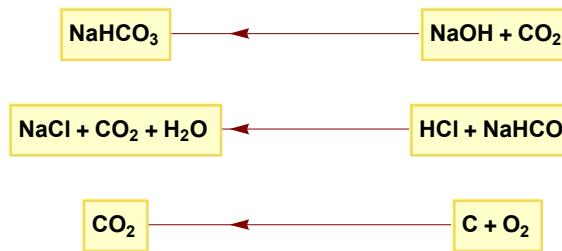
# Graphs of reactions

## Petri (13)

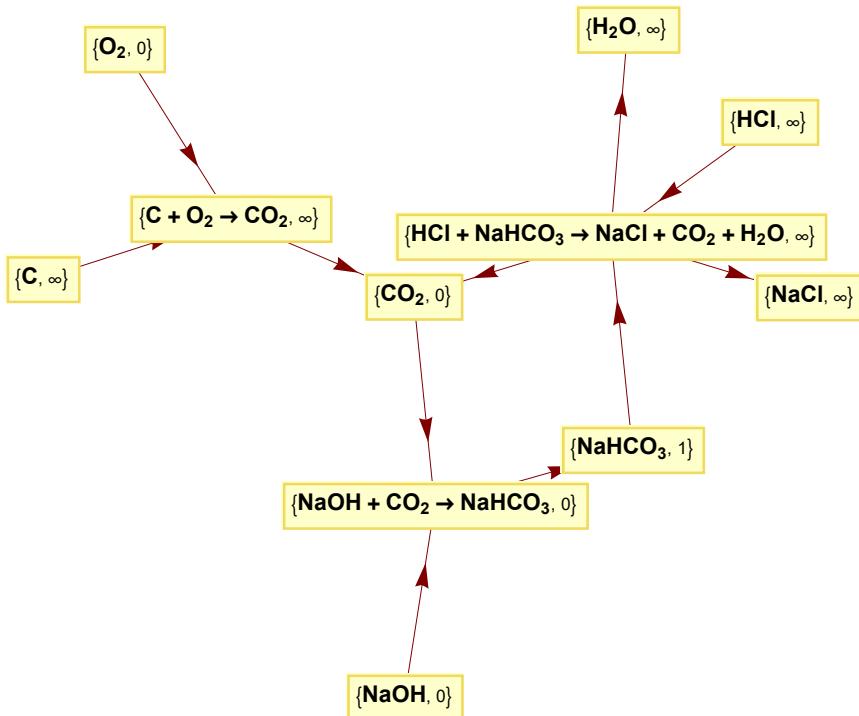
```
sm = ReactionsData[{"Petri"}][{"species", "complexes", "reactionsteps"}]
```

```
{ {C, O2, CO2, NaOH, NaHCO3, HCl, NaCl, H2O},
  {C + O2, CO2, NaOH + CO2, NaHCO3, HCl + NaHCO3, NaCl + CO2 + H2O},
  {C + O2 → CO2, NaOH + CO2 → NaHCO3, HCl + NaHCO3 → NaCl + CO2 + H2O} }
```

```
ShowFHJGraph[{"Petri"}, DirectedEdges → True, VertexLabeling → All,
  ImageSize → 300, Method → "SpringEmbedding"] /. format[Flatten[sm], Bold, 12]
```



```
ShowVolpertGraph[{"Petri"}, DirectedEdges → True,
  VertexLabeling → All, ImageSize → 500, Method → "SpringEmbedding",
  Indexed → {"CO2", "NaOH", "O2"}] /. format[Flatten[sm], Bold, 12]
```

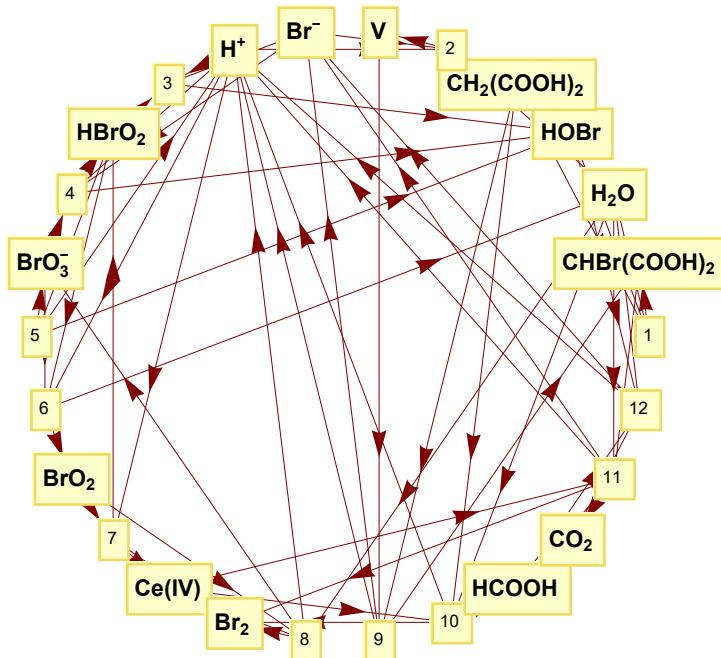


## Clarke

```

alias = {"A" → "CH2(COOH)2", "B" → "BrO3-", "H" → "H+", "M" → "CHBr(COOH)2", "P" → "HCOOH", "Q" → "CO2", "S" → "H2O", "T" → "HOBr", "U" → "BrO2", "W" → "Br2", "X" → "HBrO2", "Y" → "Br-", "Z" → "Ce(IV)"};
speccomp = Flatten[ReactionsData[{"Clarke"}][{"species", "complexes"}]] /. alias;
clarkeV = ShowVolpertGraph[GetReaction[{"Clarke"}]] /. alias, DirectedEdges → True,
VertexLabeling → True, EdgeLabeling → False, ImageSize → 400,
Method → "CircularEmbedding", Numbered → True] /. format[speccomp, Bold, 12]

```



## Acyclicity of the Volpert graph

has similar consequences as weak reversibility and zero deficiency together.

```
AcyclicGraphQ[Graph[{X → Y, X + Y → Z}]]
```

True

```
AcyclicVolpertGraphQ[{X → Y, X + Y → Z}]
```

True

## Mass Conservation

$$\exists \rho > 0 : \rho^T(\beta - \alpha) = 0$$

A reaction is mass conserving if and only if  $\max_{\lambda, \rho} \{\lambda; \rho^T(\beta - \alpha) = 0, \rho \geq \lambda \cdot \mathbf{1}, \rho^T \mathbf{1} = 1\} > 0$ .

```
MassConservationRelations[{"Lotka-Volterra"}]
{ρA > 0, ρX == ρA, ρY == ρA, ρB == ρA}

MassConservationRelations[{"Lotka-Volterra"}, ExternalSpecies → {"A", "B"}]
False
```

---

Atoms conserved → mass conserved.

The methanol-formic acid esterification

```
AtomConservingQ[{ "HCOOH" + "CH3OH" → "HCOOCH3" + "H2O" }]
{True}

ToAtomMatrix[{"HCOOH", "CH3OH", "HCOOCH3", "H2O"}, FormattedOutput → True, Frame → All]
```

Atoms\Molecules	HCOOH	CH3OH	HCOOCH3	H2O
C	1	1	2	0
H	2	4	4	2
O	2	1	2	1
charge	0	0	0	0

## Decompositions

```
ElementaryReactions[{"H2", "O2", "H2O", "C", "O", "CO"}, 2]
{O2 → 2 O, H2O → H2 + O, 2 O → O2, CO → C + O, H2 + O2 → H2O + O, H2 + O → H2O, CO + H2 → C + H2O,
 C + O2 → CO + O, C + H2O → CO + H2, H2O + O → H2 + O2, C + O → CO, CO + O → C + O2}

Decompositions[{ "H" + "O2" + 3 "H2" → 3 "H" + 2 "H2O" },
 { "H" + "O2" → "O" + "OH", "O" + "H2" → "H" + "OH", "OH" + "H2" → "H" + "H2O",
  2 "H" → "H2", "H" + "OH" ⇌ "H2O"}, 10, Method → LPBased]

7 solution(s) found with 10 solved LPs.

{{1, 1, 2, 0, 0, 0}, {1, 1, 3, 1, 0, 1}, {1, 1, 2, 0, 1, 1},
 {1, 1, 4, 2, 0, 2}, {1, 1, 3, 1, 1, 2}, {1, 1, 2, 0, 2, 2}, {1, 1, 2, 0, 3, 3}}
```

What is a kinetic differential equation?

```
DeterministicModel[{FHIII}, {k1, k-1, k2, k3, k4}, {x, y, z, t, u}][[1]] /. {w_[t] → w} // Column
x' = y k-1 - x k1 - x z k2 + u y k4
y' = -y k-1 + x k1 + t k3 - u y k4
z' = -x z k2 + u y k4
t' = x z k2 - t k3
u' = t k3 - u y k4

Σm=1 α(m, r) Xm → Σm=1 β(m, r) Xm (r = 1, 2, ..., R)
```

$$\dot{c}_m = \sum_{r=1}^R (\beta(m, r) - \alpha(m, r)) k_r \prod_{p=1}^M c_m^{\alpha(p, r)}, \text{ or shortly}$$

$$\dot{c} = \gamma \cdot \text{diag}(k) \cdot c^\alpha = P \circ c$$

## ■ What it is not?

### Generalized Lotka-Volterra model

$$\dot{x}_m = x_m \left( \sum_{p=1}^M a_{mp} x_p + b_m \right)$$

```
DeterministicModel[{3 "X" → 4 "X"}, {k}] [[1]]
```

$$\{c_X'[t] == k c_X[t]^3\}$$

```
DeterministicModel[{"Robertson"}, {k1, k2, k3}, {a, b, c}] [[1]] // Column
```

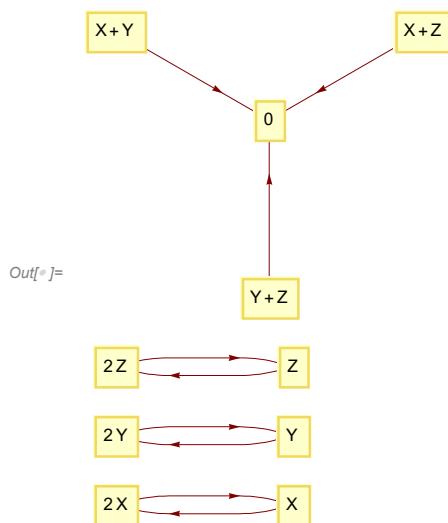
$$a'[t] == -a[t] k_1 + b[t] c[t] k_3$$

$$b'[t] == a[t] k_1 - b[t]^2 k_2 - b[t] c[t] k_3$$

$$c'[t] == b[t]^2 k_2$$

But: All the LV models are kinetic differential equations.

```
In[6]:= DeterministicModel[(antonov =
    {"X" ↔ 2 "X", "Y" ↔ 2 "Y", "Z" ↔ 2 "Z", "X" + "Y" → 0, "Y" + "Z" → 0, "Z" + "X" → 0}), {1, 1, 1, 1, 1, 1, k1, k2, k3}, {x, y, z}] [[1]] // Column
x'[t] == x[t] - x[t]^2 - k1 x[t] y[t] - k3 x[t] z[t]
Out[6]= y'[t] == y[t] - k1 x[t] y[t] - y[t]^2 - k2 y[t] z[t]
z'[t] == z[t] - k3 x[t] z[t] - k2 y[t] z[t] - z[t]^2
In[7]:= ShowFHJGraph[{antonov}, DirectedEdges → True, VertexLabeling → All, ImageSize → 200]
```



Valery Antonov, Diana Dolicanin, Valery G. Romanovski, János Tóth: Invariant Planes and Periodic Oscillations in the May-Leonard Asymmetric Model, *MATCH Commun. Math. Comput. Chem.* **76** (2016) 455-474.

## Kolmogorov systems

$$\dot{x}_m = x_m f_m \circ \mathbf{x}$$

See above, or

```
DeterministicModel[{"Brusselator"}, {a, 1, k1, b},
  {x, y}, ExternalSpecies → {"A", "B", "D", "E"}][[1]] // Column
x'[t] == a - x[t] - b x[t] + k1 x[t]^2 y[t]
y'[t] == x[t] - k1 x[t]^2 y[t]
```

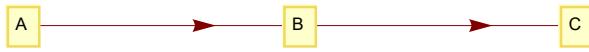
However, here is a Kolmogorov system

```
DeterministicModel[
  {2 "X" + "Y" → 3 "X" + "Y", "X" + 2 "Y" → "X" + 3 "Y"}, {k1, k2}, {x, y}][[1]] // Column
x'[t] == k1 x[t]^2 y[t]
y'[t] == k2 x[t] y[t]^2
```

## Monotone systems

$$\dot{\mathbf{x}} = \mathbf{f} \circ \mathbf{x} \quad \mathbf{y}(0) \leq \mathbf{z}(0) \text{ implies } \mathbf{y}(t) \leq \mathbf{z}(t)$$

```
ShowFHJGraph[{"Consecutive"}]
```



More generally (De Leenheer, Angeli, Sontag):

De Leenheer P., Angeli D., Sontag E. D. Monotone chemical reaction networks. *J. Math. Chem.* **41**(3) (2006), 295–314.

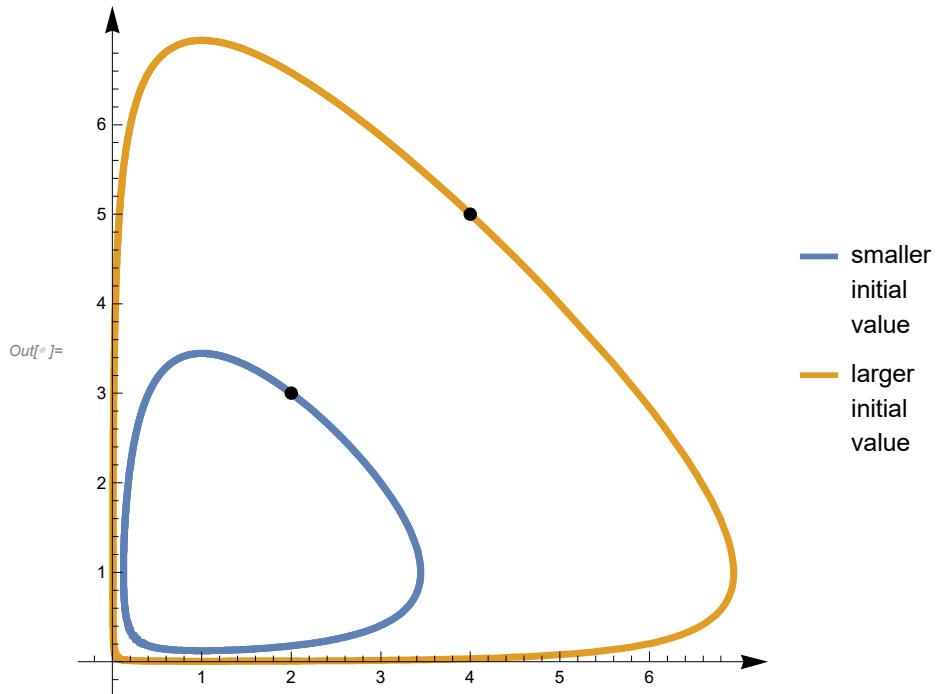
$\mathcal{C}_1 \leftrightarrow \mathcal{C}_2 \leftrightarrow \dots \leftrightarrow \mathcal{C}_{N-1} \leftrightarrow \mathcal{C}_N$  (no common species in the complexes)

But in general, induced kinetic differential equations are not monotone systems.

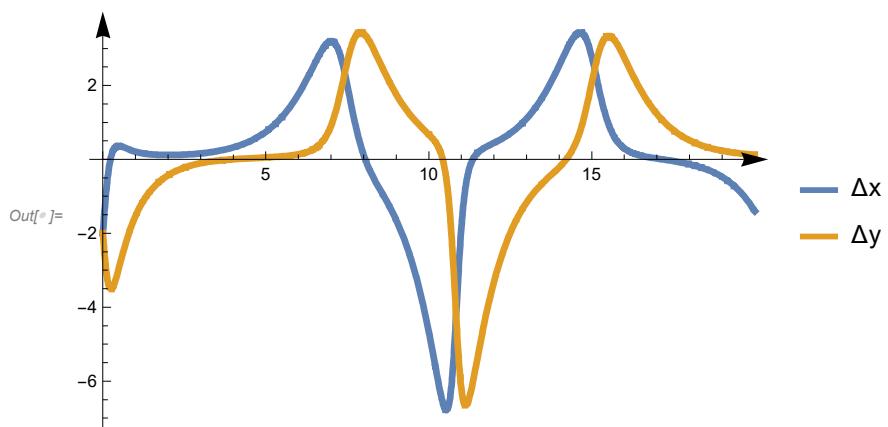
```
In[6]:= re = ReactionsData[{"Lotka-Volterra"}, ExternalSpecies → {"A", "B"}][{"fhjgraphedges"}]
Out[6]= {X → 2 X, X + Y → 2 Y, Y → 0}

In[7]:= small = ReplaceAll @@ Concentrations[{re}, {1, 1, 1}, {2, 3}, {0, 20}, {x, y}];
In[8]:= large = ReplaceAll @@ Concentrations[{re}, {1, 1, 1}, {4, 5}, {0, 20}, {x, y}];
```

```
In[6]:= lvtwocurves1 = ParametricPlot[Evaluate[{small, large}],
{t, 0, 20}, PlotRange -> All, PlotStyle -> Directive[Thickness[0.01]],
PlotLegends -> {"smaller\ninitial\nvalue", "larger\ninitial\nvalue"},
Epilog -> {PointSize[Large], Point[{2, 3}], Point[{4, 5}]}]
```



```
In[7]:= lvtwocurves2 = Plot[Evaluate[{small - large}], {t, 0, 20}, PlotRange -> All,
PlotStyle -> Directive[Thickness[0.01]], PlotLegends -> {" $\Delta x$ ", " $\Delta y$ "}]
```



## ■ What it is?

# A polynomial system without negative cross effect (Hárs-Tóth, 1979).

```

Column[{x' == y, y' == Framed[-x]}]
x' == y
y' == -x

Column[{x' == σy - σx, y' == ρx - Framed[-xz], z' == xy - βz}]
x' == -x σ + y σ
y' == x ρ - -xz
z' == xy - z β

In[°]:= CrossEffectQ[polyval_, vars_] := Module[{M = Length[vars]},
  And @@ (Map[Not[Negative[#]] &, Flatten[MapThread[ReplaceAll,
    {MonomialList[polyval, vars],
     Thread[vars → #] & /@ (1 - IdentityMatrix[M])}]]])
]

In[°]:= CrossEffectQ[{y, -x}, {x, y}]
Out[°]= False

In[°]:= RightHandSide[{FHIII}, {k1, k-1, k2, k3, k4}, {x, y, z, t, u}]
Out[°]= {y k-1 - x k1 - x z k2 + u y k4, -y k-1 + x k1 + t k3 - u y k4, -x z k2 + u y k4, x z k2 - t k3, t k3 - u y k4}

```

```

In[°]:= CrossEffectQ[
  RightHandSide[{FHIII}, {k1, k-1, k2, k3, k4}, {x, y, z, t, u}], {x, y, z, t, u}]
Out[°]= !Negative[k4] && !Negative[k-1] && !Negative[k1] &&
  !Negative[k3] && !Negative[k4] && !Negative[k2] && !Negative[k3]

```

In this talk we assume **mass action** type kinetics, thus rhs of other forms are excluded—although not in the program:

```

RightHandSide[{"H2" + "Br2" → 2 "HBr"}, {k "H2" "Br2"^(3/2) / (K "HBr" + "Br2")}, MassAction → False]
{
$$-\frac{k c_{Br2}^{3/2} c_{H2}}{c_{Br2} + K c_{HBr}}, -\frac{k c_{Br2}^{3/2} c_{H2}}{c_{Br2} + K c_{HBr}}, \frac{2 k c_{Br2}^{3/2} c_{H2}}{c_{Br2} + K c_{HBr}}}$$
}

```

What is a polynomial? Homework:  $f_m(\mathbf{x}) = \sum a_k^m \mathbf{x}^k$  if and only if the coordinate functions of  $\mathbf{f}$  are univariate polynomials of all the variables. (Carroll, 1961.)

## Equivalent forms

$$\dot{\mathbf{c}} = \gamma \cdot \text{diag}(\mathbf{k}) \cdot \mathbf{c}^\alpha = \mathbf{P} \circ \mathbf{c}$$

## Network structure emphasized

$$\dot{c} = Y \cdot E \cdot \text{diag}(k) \cdot c^\alpha \quad (E \text{ is the vertex-edge incidence matrix of the reaction})$$

Lotka-Volterra:

```
ReactionsData[{"Lotka-Volterra"}, ExternalSpecies → {"A", "B"}] ["complexes"]
{X, 2 X, X + Y, 2 Y, Y, 0}
```

```
α = ReactionsData[{"Lotka-Volterra"}, ExternalSpecies → {"A", "B"}] ["α"]
```

SparseArray [  Specified elements: 4 ]  
Dimensions: {2, 3}

$$Y = \begin{pmatrix} 1 & 2 & 1 & 0 & 0 & 0 \\ 0 & 0 & 1 & 2 & 1 & 0 \end{pmatrix}; \quad E = \begin{pmatrix} -1 & 0 & 0 \\ 1 & 0 & 0 \\ 0 & -1 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & -1 \\ 0 & 0 & 1 \end{pmatrix}; \quad kk = \begin{pmatrix} k_1 \\ k_2 \\ k_3 \end{pmatrix};$$

```
Y.E.(kk Times @@ {x, y}^α) // Flatten
{x k₁ - x y k₂, x y k₂ - y k₃}
```

```
RightHandSide[{"Lotka-Volterra"}, {k₁, k₂, k₃}, {x, y}, ExternalSpecies → {"A", "B"}]
{x k₁ - x y k₂, x y k₂ - y k₃}
```

How to find the incidence matrix? Use the function IncidenceMatrix ⊕

```
lot = Graph[ReactionsData[{"Lotka-Volterra"}] ["fhjgraphedges"]];
TableForm[IncidenceMatrix[#, 
  TableHeadings → {VertexList[#, EdgeList[#]], TableAlignments → Right} &[lot]]]
```

	A + X ↔ 2 X	X + Y ↔ 2 Y	Y ↔ B
A + X	-1	0	0
2 X	1	0	0
X + Y	0	-1	0
2 Y	0	1	0
Y	0	0	-1
B	0	0	1

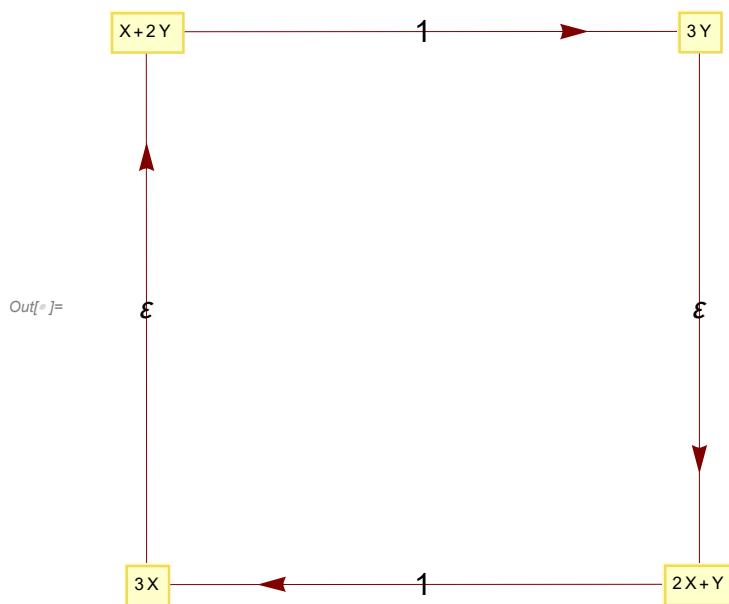
## Linearity in the reaction rate coefficients emphasized

$$\dot{c} = y \cdot \text{diag}(c^\alpha) \cdot k$$

# Stationary points

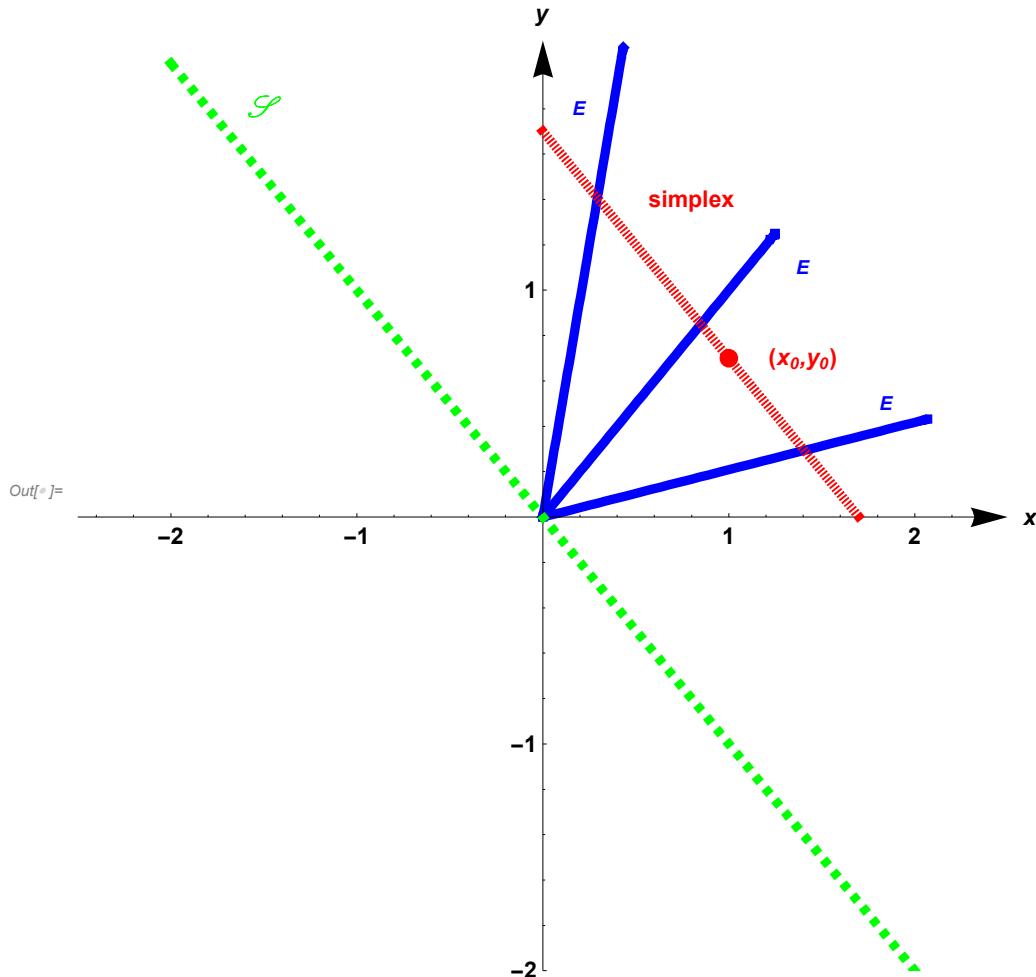
## ■ Existence, uniqueness

```
In[®]:= hjreaction = ShowFHJGraph[
  {2 "X" + "Y" → 3 "X", 3 "X" → "X" + 2 "Y", "X" + 2 "Y" → 3 "Y", 3 "Y" → 2 "X" + "Y"}, 
  {Style[1, 16], Style[ε, 16], Style[1, 16], Style[ε, 16]}, 
  DirectedEdges → True, VertexLabeling → All, Method → "SpringEmbedding"]
```



```
In[10]:= ClearAll[statcurves, m, n, u, a, b, ε, x, y];
statcurves[u_] := PowerExpand[{x, y} /. Cases[
  Last[Simplify[StationaryPoints[{"Horn-Jackson"}, {ε, 1, ε, 1}, {m, n}, {x, y}], Assumptions → (0 < ε < 1/6)]], {a_?TrueQ, b___} → {b}] /. m → (u - n) /. ε → 1/12]

hornjackson = ParametricPlot[Evaluate[statcurves[u]], {u, 0, 2.5},
  PlotStyle → Directive[{Blue, Thickness[0.01]}], AspectRatio → 1,
  Epilog → {Dashed, Green, Thickness[0.01], Line[{{-2, 2}, {2, -2}}], Dashing[{0}],
    Red, Thickness[0.01], Line[{{0, 1.7}, {1.7, 0}}], PointSize[0.02],
    Point[{1.0, 0.7}], Text[Style["(x0,y0)", Italic, Bold, 12], {1.4, 0.7}],
    Green, Text[Style["S", FontFamily → "Kunstler Script", Bold, 16], {-1.5, 1.8}],
    Red, Text[Style["simplex", Bold, 12], {0.8, 1.4}], Blue,
    Text[Style["E", Italic, Bold, 10], {1.85, 0.5}], Text[Style["E", Italic, Bold, 10], {1.4, 1.1}],
    Text[Style["E", Italic, Bold, 10], {0.2, 1.8}]},
  PlotRange → {{-2.5, 2.5}, {-2.0, 2.1}}, AxesLabel →
  (Style[#, Bold, Italic, 12] & /@ {"x", "y"}),
  TicksStyle → Directive[Black, Bold, 12], ImageSize → 500]
```



Weak reversibility does not imply uniqueness. However,

Weak reversibility implies existence of stationary points in all positive stoichiometric compatibility classes (Boros, 2017).

## ■ Absolute Concentration Robustness

```
StationaryPoints[{"X" + "Y" → 2 "Y", "Y" → "X"}, {k1, k2}, {x0, y0}][[2, 1]]
{ $k_1 > \frac{k_2}{x_0 + y_0}$ ,  $c_X^* \rightarrow \frac{k_2}{k_1}$ ,  $c_Y^* \rightarrow \frac{-k_2 + k_1 x_0 + k_1 y_0}{k_1}$ }
```

Now matter what the stationary concentration is, the first component does not depend on the initial concentrations: A kind of stability, may be useful from biological point of view. (Feinberg, Shinar).

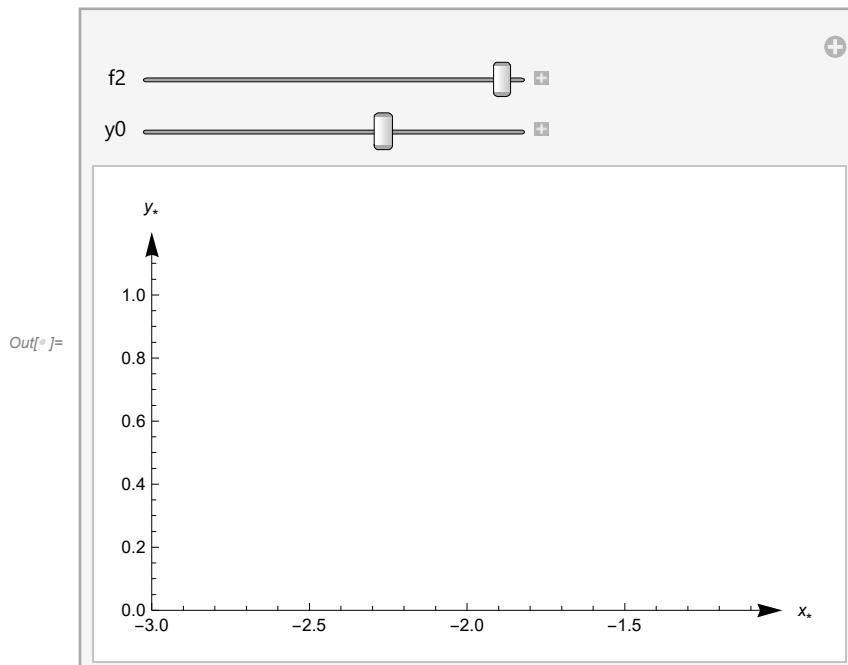
```
AbsoluteConcentrationRobustness[{"X" + "Y" → 2 "Y", "Y" → "X"}]
{X}
```

Also in the stochastic model.

## ■ Parameter dependence

```
x' = -x y + f (1 - x)
y' = x y + y  $\left( \left( 1 - \frac{f_2}{f} \right) y_0 - y \right) + (f - f_2) y_0 - f y$ 

In[6]:= liliipota[f2_, y0_, f_] := f x3 - f x2 + x (f2 + (f - f2) y0) - f2;
Manipulate[
ParametricPlot[(XX[f2_, y0_, j_] := x /. NSolve[liliipota[f2, y0, 10j] == 0, {x}]];
Distribute[{j, XX[f2, y0, j]}, List]), {j, -3, -1},
PlotRange → {{-3, -1}, {0, 1.2}}, AxesLabel → {"x*", "y*"}],
{{f2, 1 / 135}, 0.0009, 0.0075, 0.0001}, {{y0, 10 / 27}, 0.24, 0.38, 0.01}]
```



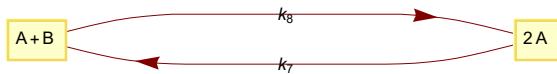
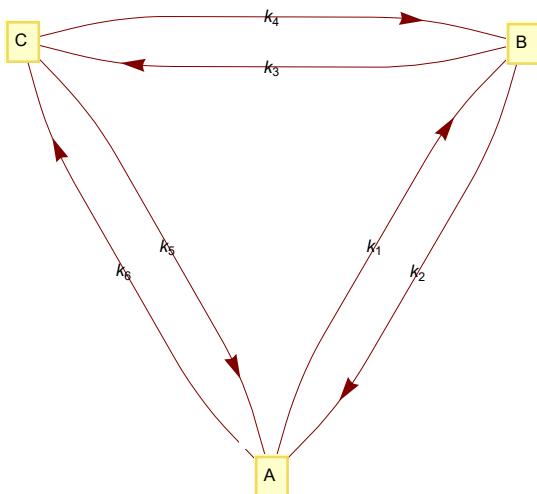
## ■ Detailed balance

A reaction only consisting of reversible steps is detailed balanced at the positive stationary concentration  $\mathbf{c}_*$ , if for all steps  $k_p \mathbf{c}_*^{\alpha(.,p)} = k_{-p} \mathbf{c}_*^{\beta(.,p)}$  holds. (Conditions: Feinberg; Fredholm)

Nagy, I., Tóth, J.: Microscopic reversibility or detailed balance in ion channel models. *J. Math. Chem.* **50** (5) (2012), 1179–1199.

Nagy, I., Kovács, B., Tóth, J.: Detailed balance in ion channels: Applications of Feinberg's theorem. *React. Kinet. Catal. Lett.* **96** (2) (2009), 263–267.

```
In[®] = ShowFHJGraph[
  complicated = Join[GetReaction[{"Wegscheider"}], ToReversible["Triangle"]],
  {k1, k2, k7, k8, k3, k4, k5, k6}, EdgeLabeled → True]
```



```
DetailedBalanced[complicated, {k1, k2, k7, k8, k3, k4, k5, k6}]
```

$$\{k_2 k_4 k_6 = k_1 k_3 k_5, k_2 k_7 = k_1 k_8\}$$

Tóth, J.; Nagy, A. L.; Zsély, I. Gy.: Structural analysis of combustion mechanisms, *J. Math. Chem.* **53** (2015), 86–110.

In a mechanism for hydrogen combustion (9 species, 7 independent reaction steps,  $\delta = 41 - 16 - 7 = 18$ , ) given by Starik et al. (2009)

A.M. Starik, N.S. Titova, A.S. Sharipov, V.E. Kozlov: Syngas oxidation mechanism. *Combust. Explos. Shock Waves* **46**(5) (2010), 491–506.

$$\begin{aligned} \{ & k_2 k_4 k_5 = k_1 k_3 k_6, k_3 k_8 k_{11} = k_4 k_7 k_{12}, k_2 k_{14} k_{19} = k_1 k_{13} k_{20}, \\ & k_4 k_{28} k_{31} = k_3 k_{27} k_{32}, k_4 k_{13} k_{41} = k_3 k_{14} k_{42}, k_{18} k_{21} k_{32} = k_{17} k_{22} k_{31}, \\ & k_2 k_9 k_{14} k_{15} = k_1 k_{10} k_{13} k_{16}, k_2 k_8 k_9 k_{21} = k_1 k_7 k_{10} k_{22}, k_2 k_4 k_9 k_{23} = k_1 k_3 k_{10} k_{24}, \\ & k_2 k_4 k_{30} k_{31} = k_1 k_3 k_{29} k_{32}, k_3 k_{14} k_{32} k_{35} = k_4 k_{13} k_{31} k_{36}, k_1 k_3 k_{10} k_{32} k_{33} = k_2 k_4 k_9 k_{31} k_{34}, \\ & k_3 k_{14} k_{32} k_{38} k_{43} = k_4 k_{13} k_{31} k_{37} k_{44}, k_2 k_4^2 k_8 k_9 k_{13} k_{26} k_{31}^2 = k_1 k_3^2 k_7 k_{10} k_{14} k_{25} k_{32}^2, \\ & k_2 k_4 k_8 k_9 k_{13} k_{31} k_{39} = k_1 k_3 k_7 k_{10} k_{14} k_{32} k_{40}, k_4 k_8 k_9 k_{13}^2 k_{31} k_{37} k_{49} = k_3 k_7 k_{10} k_{14}^2 k_{32} k_{38} k_{50}, \\ & k_2 k_4 k_8 k_9 k_{31} k_{37} k_{51} = k_1 k_3 k_7 k_{10} k_{32} k_{38} k_{52}, k_1 k_3^2 k_7 k_{10} k_{14} k_{32} k_{38} k_{45} = k_2 k_4^2 k_8 k_9 k_{13} k_{31}^2 k_{37} k_{46}, \\ & k_2 k_4 k_8 k_9 k_{13} k_{31} k_{37} k_{47} = k_1 k_3 k_7 k_{10} k_{14} k_{32} k_{38} k_{48} \} \end{aligned}$$

Carbon monoxide, methanol: more complicated.

## ■ First integrals

Nagy, I., Tóth, J.: Quadratic first integrals of kinetic differential equations. *J. Math. Chem.* **52**(1) (2014), 93–114.

# Numerics

## ■ Stiff equations

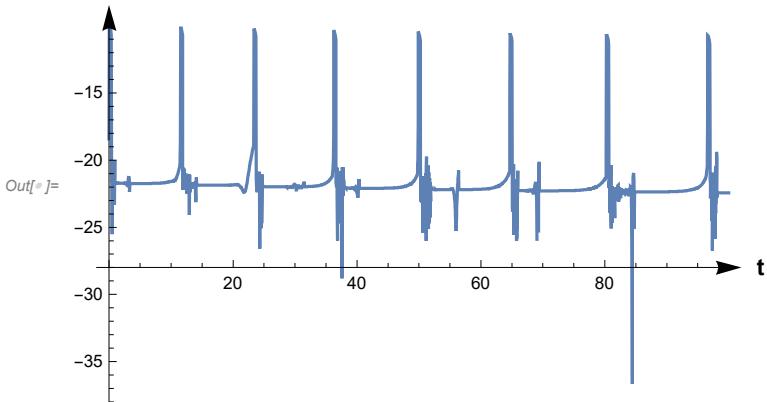
Taken from the book by Deuflhard and Bornemann who give the reaction rate coefficients (pp. 17–18), but no concentrations, still one can reproduce their results quite well.

### Oregonator - the role of options

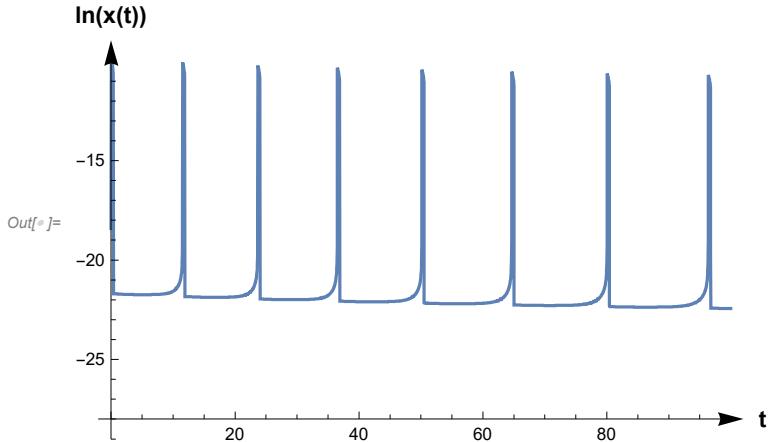
```
In[1]:= ClearAll[a, y, x, p];
or = GetReaction["Oregonator"] /. {fOregonator → 1} /. {"B" → "A", "Q" → "P"}
Out[1]= {A + Y → X, X + Y → P, A + X → 2 X + Z, 2 X → P, Z → Y}

In[2]:= cin = Concentrations[{or}, {134 / 100, 16 × 108, 8 × 103, 4 × 107, 1},
{5 × 10-1, 10-6, 10-8, 6 × 10-2, 10-4}, {0, 100}, {a, y, x, p, z}];
bzdeuflhardA = Plot[Log[x[t] /. Last[cin]], {t, 0, 100}, PlotRange → All,
AxesLabel → (Style[#, Bold, 12] & /@ {"t", "ln(x(t))"}), AxesOrigin → {0, -28}]
```

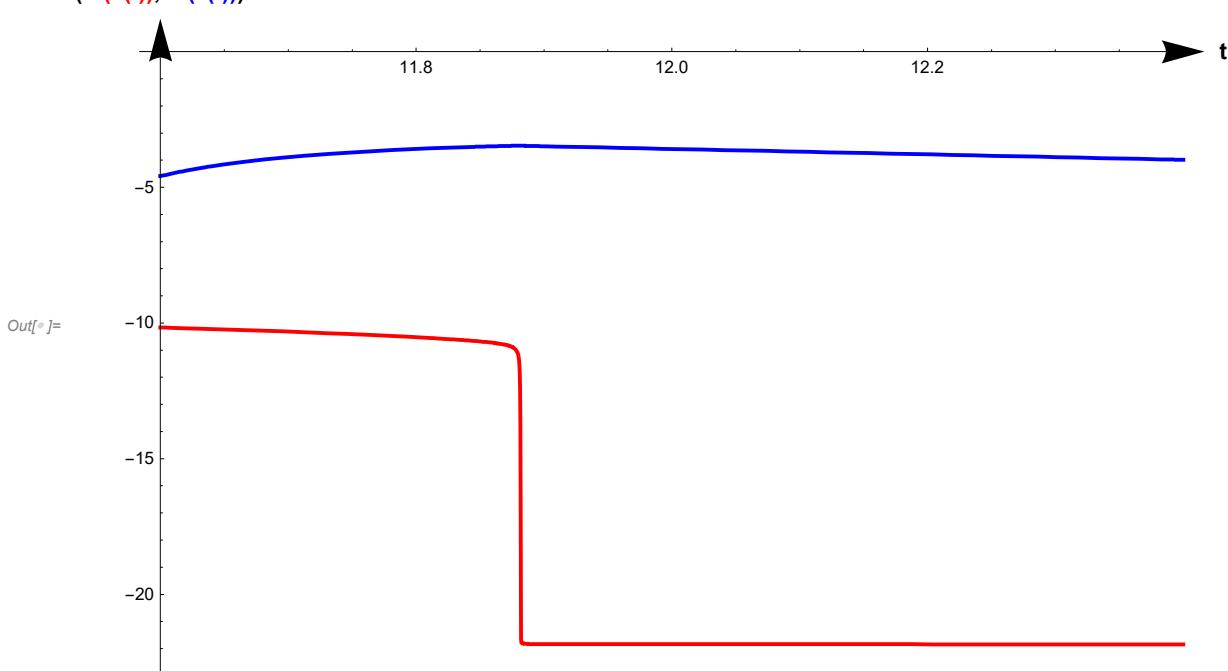
ln(x(t))



```
In[6]:= con = Concentrations[{or}, {134 / 100, 16 × 108, 8 × 103, 4 × 107, 1},
  {5 × 10-1, 10-6, 10-8, 6 × 10-2, 10-4}, {0, 100}, {a, y, x, p, z},
  Method → "BDF", WorkingPrecision → 32, MaxSteps → 106];
bzdeuflhardB = Plot[Log[x[t] /. Last[con]], {t, 0, 100}, PlotRange → All,
  AxesLabel → (Style[#, Bold, 12] & /@ {"t", "ln(x(t))"}), AxesOrigin → {0, -28}]
```



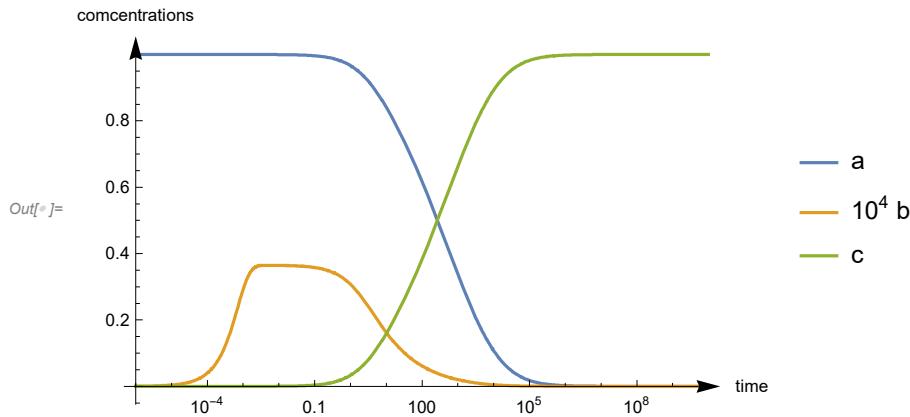
```
In[7]:= bzdeuflhardStiff = Plot[Evaluate[Log[{x[t], z[t]} /. Last[con]]],
  {t, 11.6, 12.4}, PlotStyle → {Directive[Thick, Red], Directive[Thick, Blue]},
  AxesLabel → {Style["t", Bold, 12],
  Row[{Style["(", Bold, 12], Style["ln(x(t))", Red, Bold, 12], Style[", ", Bold, 12],
  Style["ln(z(t))", Blue, Bold, 12], Style[")", Bold, 12]}]}, ImageSize → 600]
```



## Robertson - as ODE, as DAE

### ODE

```
In[1]:= conrob = Concentrations[{rob = GetReaction["Robertson"]}, {0.04, 3×107, 104}, {1, 0, 0}, {0, 1010}];
robertsonexact = LogLinearPlot[Evaluate[{1, 104, 1} (ReplaceAll @@ conrob)], {t, 10-6, 1010}, PlotLegends → {"a", "104 b", "c"}, PlotRange → All, AxesLabel → {"time", "concentrations"}]
```



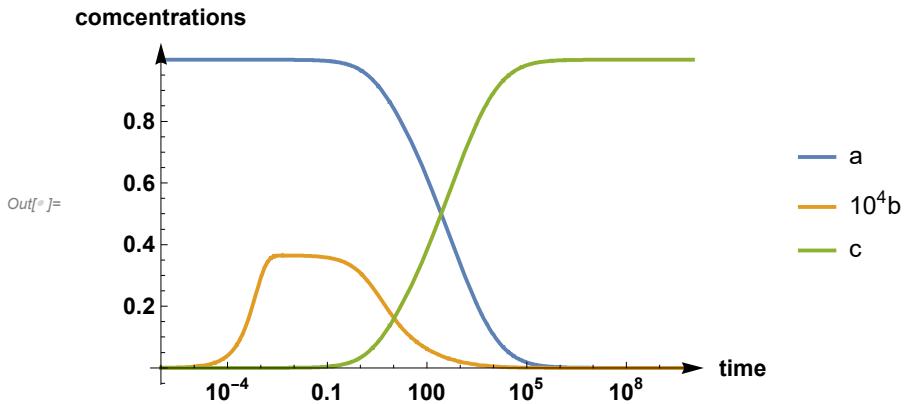
Why is it interesting (“a challenge”)?

```
dm = DeterministicModel[{rob}, {0.04, 3×107, 104}, {a, b, c}]
{{a'[t] == -0.04 a[t] + 10000 b[t] c[t],
b'[t] == 0.04 a[t] - 30000000 b[t]2 - 10000 b[t] c[t],
c'[t] == 30000000 b[t]2}, {a[t], b[t], c[t]}}
```

### DAE

```
In[2]:= nds = NDSolve[{a'[t] == -0.04 a[t] + 10000 b[t] c[t], a[t] + b[t] + c[t] == 1,
c'[t] == 30000000 b[t]2, a[0] == 1, c[0] == 0}, {a[t], b[t], c[t]}, {t, 10-6, 1010}];
```

```
In[8]:= robertsonDAE = LogLinearPlot[Evaluate[{1, 104, 1}{a[t], b[t], c[t]} /. nds], {t, 10-6, 1010}, PlotLegends -> {"a", "104b", "c"}, PlotRange -> All, PlotStyle -> Thick, AxesLabel -> (Style[#, 12, Bold] & /@ {"time", "concentrations"}), TicksStyle -> Directive[Black, Bold, 12]]
```



Quite similar to the exact figure.

## ■ Can you find the stationary point?

```
rhs = RightHandSide[{0 → X1 ↔ X2 ↔ X3 ↔ X4 ↔ X5 ↔ X6 ↔ X7 ↔ X8 → 0}, Join[{k0}, Flatten[Transpose[{k# & /@ Range[7], k# & /@ -Range[7]}]], {k8}], x# & /@ Range[8]]
```

We can only verify it:

```
rhs /. {xi_ → k0 Sum[Product[k1, {l, 10 - j, 8}], Product[k-1, {l, i, 8 - j}] / Product[k1, {l, i, 8}], {j, 1, 9 - i}]} // Simplify
```

It works in simple cases :)

```
StationaryPoints[{2 "X" ↔ "Y"}, {x, λ}, {x0, y0}] // PowerExpand
```

```
Out[8]= {{cX*, cY*}, {{cX* → -1/(4 K) (λ - √λ) √(8 x0 K + 16 y0 K + λ), cY* → 1/(8 K) (4 x0 K + 8 y0 K + λ - √λ) √(8 x0 K + 16 y0 K + λ)}}}}
```

## ■ Good numerical methods

The Euler method when applied with a small enough step size keeps the invariance of the first orthant.

The stationary points of an autonomous differential equation are the same as those obtained by the Euler method.

# Stochastic model

## ■ Symbolic calculations (limited:)

```
In[1]:= gene = {"inactive gene" ↔ "active gene" → "messenger" → "protein" → 0, "messenger" → 0};

In[2]:= ClearAll[i, a, m, p, G];
MasterEquation[{gene}, {λ1, λ-1, λ2, λ3, γ+, γ-}, {t, i, a, m, p}, P]

Out[2]= {P(1,0,0,0,0)[t, i, a, m, p] == (1 + m) P[t, i, a, 1 + m, p] γ- +
(1 + p) P[t, i, a, m, 1 + p] γ+ + (1 + a) P[t, -1 + i, 1 + a, m, p] λ-1 +
(1 + i) P[t, 1 + i, -1 + a, m, p] λ1 + (1 + a) P[t, i, 1 + a, -1 + m, p] λ2 +
(1 + m) P[t, i, a, 1 + m, -1 + p] λ3 - P[t, i, a, m, p] (m γ- + p γ+ + a λ-1 + i λ1 + a λ2 + m λ3)}

In[3]:= ProbabilityGeneratingFunctionEquation[
{gene}, {λ1, λ-1, λ2, λ3, γ+, γ-}, {t, x, y, u, v}, G]

Out[3]= G(1,0,0,0,0)[t, x, y, u, v] ==
(1 - v) γ+ G(0,0,0,0,1)[t, x, y, u, v] + (1 - u) γ- G(0,0,0,1,0)[t, x, y, u, v] +
(-u + v) λ3 G(0,0,0,1,0)[t, x, y, u, v] + (x - y) λ-1 G(0,0,1,0,0)[t, x, y, u, v] +
(u - y) λ2 G(0,0,1,0,0)[t, x, y, u, v] + (-x + y) λ1 G(0,1,0,0,0)[t, x, y, u, v]

In[4]:= SolveProbabilityGeneratingFunctionEquation[{gene}, {λ1, λ-1, λ2, λ3, γ+, γ-},
{1, 0, 0, 0}, {t, x, y, u, v}, G, Method → "MatrixExponential"]

... SolveProbabilityGeneratingFunctionEquation: The order of the reaction is at most one, thus the method "Characteristics" can also be applied.

Out[4]= {G[t, x, y, u, v],
```

$$\begin{aligned} G[t, x, y, u, v] \rightarrow 1 + (-1 + u) \left( \left( e^{t(-\gamma_- - \lambda_3)} \lambda_2 (\gamma_- \lambda_1 - \gamma_+ \lambda_1 + \lambda_1 \lambda_3) \right) / ((\gamma_- - \gamma_+ + \lambda_3) \right. \\ \left. (\gamma_-^2 - \gamma_- \lambda_{-1} - \gamma_- \lambda_1 - \gamma_- \lambda_2 + \lambda_1 \lambda_2 + 2 \gamma_- \lambda_3 - \lambda_{-1} \lambda_3 - \lambda_1 \lambda_3 - \lambda_2 \lambda_3 + \lambda_3^2) \right) - \\ \left( e^{\frac{1}{2} t (-\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2})} \lambda_2 \left( -2 \gamma_+ \lambda_1 + \lambda_{-1} \lambda_1 + \lambda_1^2 + \lambda_1 \lambda_2 + \right. \right. \\ \left. \left. \lambda_1 \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) / \left( 2 \left( \gamma_- \gamma_+ \lambda_{-1} + \gamma_- \gamma_+ \lambda_1 + \gamma_- \gamma_+ \lambda_2 + \gamma_- \lambda_1 \lambda_2 + \right. \right. \\ \left. \left. \gamma_+ \lambda_1 \lambda_2 + \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^3 + \gamma_+ \lambda_{-1} \lambda_3 + \right. \right. \\ \left. \left. \gamma_+ \lambda_1 \lambda_3 + \gamma_+ \lambda_2 \lambda_3 + \lambda_1 \lambda_2 \lambda_3 + \frac{1}{4} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^2 \right. \right. \\ \left. \left. (3 \gamma_- + 3 \gamma_+ + 3 \lambda_{-1} + 3 \lambda_1 + 3 \lambda_2 + 3 \lambda_3) + \right. \right. \\ \left. \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) (2 \gamma_- \gamma_+ + 2 \gamma_- \lambda_{-1} + 2 \gamma_+ \lambda_{-1} + 2 \right. \right. \\ \left. \left. \gamma_- \lambda_1 + 2 \gamma_+ \lambda_1 + 2 \gamma_- \lambda_2 + 2 \gamma_+ \lambda_2 + 2 \lambda_1 \lambda_2 + 2 \gamma_+ \lambda_3 + 2 \lambda_{-1} \lambda_3 + 2 \lambda_1 \lambda_3 + 2 \lambda_2 \lambda_3) \right) \right) - \\ \left( e^{\frac{1}{2} t (-\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2})} \lambda_2 \left( -2 \gamma_+ \lambda_1 + \lambda_{-1} \lambda_1 + \lambda_1^2 + \lambda_1 \lambda_2 - \right. \right. \\ \left. \left. \lambda_1 \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) \end{aligned}$$

$$\begin{aligned}
& \left. \left( \lambda_1 \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) / \left( 2 \left( \gamma_- \gamma_+ \lambda_{-1} + \gamma_- \gamma_+ \lambda_1 + \gamma_- \gamma_+ \lambda_2 + \gamma_- \lambda_1 \lambda_2 + \right. \right. \\
& \gamma_+ \lambda_1 \lambda_2 + \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^3 + \gamma_+ \lambda_{-1} \lambda_3 + \\
& \gamma_+ \lambda_1 \lambda_3 + \gamma_+ \lambda_2 \lambda_3 + \lambda_1 \lambda_2 \lambda_3 + \frac{1}{4} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^2 \\
& (3 \gamma_- + 3 \gamma_+ + 3 \lambda_{-1} + 3 \lambda_1 + 3 \lambda_2 + 3 \lambda_3) + \\
& \left. \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) (2 \gamma_- \gamma_+ + 2 \gamma_- \lambda_{-1} + 2 \gamma_+ \lambda_{-1} + 2 \gamma_- \right. \right. \\
& \left. \left. \lambda_1 + 2 \gamma_+ \lambda_1 + 2 \gamma_- \lambda_2 + 2 \gamma_+ \lambda_2 + 2 \lambda_1 \lambda_2 + 2 \gamma_+ \lambda_3 + 2 \lambda_{-1} \lambda_3 + 2 \lambda_1 \lambda_3 + 2 \lambda_2 \lambda_3) \right) \right) + \\
& (-1 + v) \left( - \left( \left( e^{-t \gamma_+} \lambda_1 \lambda_2 \lambda_3 \right) / \left( (\gamma_+^2 - \gamma_+ \lambda_{-1} - \gamma_+ \lambda_1 - \gamma_+ \lambda_2 + \lambda_1 \lambda_2) (-\gamma_- + \gamma_+ - \lambda_3) \right) \right) - \right. \\
& \left( e^{t (-\gamma_- - \lambda_3)} \lambda_1 \lambda_2 \lambda_3 \right) / \\
& \left( (\gamma_- - \gamma_+ + \lambda_3) (\gamma_-^2 - \gamma_- \lambda_{-1} - \gamma_- \lambda_1 - \gamma_- \lambda_2 + \lambda_1 \lambda_2 + 2 \gamma_- \lambda_3 - \lambda_{-1} \lambda_3 - \lambda_1 \lambda_3 - \lambda_2 \lambda_3 + \lambda_3^2) \right) + \\
& \left. \left( e^{\frac{1}{2} t (-\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2})} \lambda_1 \lambda_2 \lambda_3 \right) \right) / \\
& \left( \gamma_- \gamma_+ \lambda_{-1} + \gamma_- \gamma_+ \lambda_1 + \gamma_- \gamma_+ \lambda_2 + \gamma_- \lambda_1 \lambda_2 + \gamma_+ \lambda_1 \lambda_2 + \right. \\
& \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^3 + \gamma_+ \lambda_{-1} \lambda_3 + \gamma_+ \lambda_1 \lambda_3 + \right. \\
& \gamma_+ \lambda_2 \lambda_3 + \lambda_1 \lambda_2 \lambda_3 + \frac{1}{4} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^2 \\
& (3 \gamma_- + 3 \gamma_+ + 3 \lambda_{-1} + 3 \lambda_1 + 3 \lambda_2 + 3 \lambda_3) + \\
& \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) (2 \gamma_- \gamma_+ + 2 \gamma_- \lambda_{-1} + 2 \gamma_+ \lambda_{-1} + \right. \\
& 2 \gamma_- \lambda_1 + 2 \gamma_+ \lambda_1 + 2 \gamma_- \lambda_2 + 2 \gamma_+ \lambda_2 + 2 \lambda_1 \lambda_2 + 2 \gamma_+ \lambda_3 + 2 \lambda_{-1} \lambda_3 + 2 \lambda_1 \lambda_3 + 2 \lambda_2 \lambda_3) \right) + \\
& \left. \left( e^{\frac{1}{2} t (-\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2})} \lambda_1 \lambda_2 \lambda_3 \right) \right) / \left( \gamma_- \gamma_+ \lambda_{-1} + \gamma_- \gamma_+ \lambda_1 + \gamma_- \gamma_+ \lambda_2 + \right. \\
& \gamma_- \lambda_1 \lambda_2 + \gamma_+ \lambda_1 \lambda_2 + \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^3 + \gamma_+ \lambda_{-1} \lambda_3 + \\
& \gamma_+ \lambda_1 \lambda_3 + \gamma_+ \lambda_2 \lambda_3 + \lambda_1 \lambda_2 \lambda_3 + \frac{1}{4} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^2 \\
& (3 \gamma_- + 3 \gamma_+ + 3 \lambda_{-1} + 3 \lambda_1 + 3 \lambda_2 + 3 \lambda_3) + \\
& \left. \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) (2 \gamma_- \gamma_+ + 2 \gamma_- \lambda_{-1} + 2 \gamma_+ \lambda_{-1} + 2 \gamma_- \lambda_1 + \right. \right. \\
& 2 \gamma_+ \lambda_1 + 2 \gamma_- \lambda_2 + 2 \gamma_+ \lambda_2 + 2 \lambda_1 \lambda_2 + 2 \gamma_+ \lambda_3 + 2 \lambda_{-1} \lambda_3 + 2 \lambda_1 \lambda_3 + 2 \lambda_2 \lambda_3) \right) + (-1 + x) \\
& \left( - \left( \left( e^{t (-\gamma_- - \lambda_3)} \left( \lambda_{-1} \left( (\gamma_+ - \lambda_1) \lambda_1 + \frac{1}{2} \lambda_1 \left( -\lambda_{-1} + \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) \right) + \right. \right. \right. \\
& \left. \left. \left. \frac{1}{2} \left( \lambda_{-1} - \lambda_1 + \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right. \right. \\
& \left. \left. \left( \lambda_{-1} \lambda_1 + \frac{1}{2} (\gamma_+ - \lambda_1) \left( \lambda_{-1} - \lambda_1 + \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) \right) \right) \right) /
\end{aligned}$$

$$\begin{aligned}
& \left( (\gamma_- - \gamma_+ + \lambda_3) \left( \gamma_-^2 - \gamma_- \lambda_{-1} - \gamma_- \lambda_1 - \gamma_- \lambda_2 + \lambda_1 \lambda_2 + 2 \gamma_- \lambda_3 - \lambda_{-1} \lambda_3 - \lambda_1 \lambda_3 - \lambda_2 \lambda_3 + \lambda_3^2 \right) \right) - \\
& \left( e^{-t \gamma_+} \left( \lambda_{-1} \left( \frac{1}{2} \lambda_1 \left( -\lambda_{-1} + \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) + \right. \right. \\
& \quad \left. \frac{1}{2} \lambda_1 \left( \lambda_{-1} - \lambda_1 + \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) + \\
& \quad \left( \lambda_{-1} \lambda_1 + \frac{1}{4} \left( \lambda_{-1} - \lambda_1 + \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right. \\
& \quad \left. \left. \left. \left( \lambda_{-1} - \lambda_1 + \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) (\gamma_- - \lambda_1 + \lambda_3) \right) \Bigg) / \\
& \left( (\gamma_+^2 - \gamma_+ \lambda_{-1} - \gamma_+ \lambda_1 - \gamma_+ \lambda_2 + \lambda_1 \lambda_2) (-\gamma_- + \gamma_+ - \lambda_3) \right) + \left( e^{\frac{1}{2} t \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2} \right)} \right. \\
& \quad \left( \lambda_{-1} \left( (\gamma_+ - \lambda_1) \lambda_1 + \frac{1}{2} \lambda_1 \left( -\lambda_{-1} + \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) + \right. \\
& \quad \left. \left( \lambda_{-1} \lambda_1 + \frac{1}{2} (\gamma_+ - \lambda_1) \left( \lambda_{-1} - \lambda_1 + \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) \right. \\
& \quad \left. \left. (\gamma_- - \lambda_1 + \lambda_3) \right) \right) / \left( \gamma_- \gamma_+ \lambda_{-1} + \gamma_- \gamma_+ \lambda_1 + \gamma_- \gamma_+ \lambda_2 + \gamma_- \lambda_1 \lambda_2 + \gamma_+ \lambda_1 \lambda_2 + \right. \\
& \quad \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^3 + \gamma_+ \lambda_{-1} \lambda_3 + \gamma_+ \lambda_1 \lambda_3 + \right. \\
& \quad \left. \gamma_+ \lambda_2 \lambda_3 + \lambda_1 \lambda_2 \lambda_3 + \frac{1}{4} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^2 \right. \\
& \quad \left. (3 \gamma_- + 3 \gamma_+ + 3 \lambda_{-1} + 3 \lambda_1 + 3 \lambda_2 + 3 \lambda_3) + \right. \\
& \quad \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right. \\
& \quad \left. (2 \gamma_- \gamma_+ + 2 \gamma_- \lambda_{-1} + 2 \gamma_+ \lambda_{-1} + 2 \gamma_- \lambda_1 + 2 \gamma_+ \lambda_1 + 2 \gamma_- \lambda_2 + 2 \gamma_+ \lambda_2 + 2 \lambda_1 \lambda_2 + \right. \\
& \quad \left. 2 \gamma_+ \lambda_3 + 2 \lambda_{-1} \lambda_3 + 2 \lambda_1 \lambda_3 + 2 \lambda_2 \lambda_3) \right) + \left( e^{\frac{1}{2} t \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2} \right)} \right. \\
& \quad \left( \lambda_{-1} \left( (\gamma_+ - \lambda_1) \lambda_1 + \frac{1}{2} \lambda_1 \left( -\lambda_{-1} + \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) + \right. \\
& \quad \left. \left( \lambda_{-1} \lambda_1 + \frac{1}{2} (\gamma_+ - \lambda_1) \left( \lambda_{-1} - \lambda_1 + \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) \right. \\
& \quad \left. \left. (\gamma_- - \lambda_1 + \lambda_3) \right) \right) / \left( \gamma_- \gamma_+ \lambda_{-1} + \gamma_- \gamma_+ \lambda_1 + \gamma_- \gamma_+ \lambda_2 + \gamma_- \lambda_1 \lambda_2 + \gamma_+ \lambda_1 \lambda_2 + \right. \\
& \quad \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^3 + \gamma_+ \lambda_{-1} \lambda_3 + \gamma_+ \lambda_1 \lambda_3 + \right. \\
& \quad \left. \gamma_+ \lambda_2 \lambda_3 + \lambda_1 \lambda_2 \lambda_3 + \frac{1}{4} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right)^2 \right. \\
& \quad \left. (3 \gamma_- + 3 \gamma_+ + 3 \lambda_{-1} + 3 \lambda_1 + 3 \lambda_2 + 3 \lambda_3) + \right. \\
& \quad \left. \frac{1}{2} \left( -\lambda_{-1} - \lambda_1 - \lambda_2 + \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) (2 \gamma_- \gamma_+ + 2 \gamma_- \lambda_{-1} + 2 \gamma_+ \lambda_{-1} + \right. \\
& \quad \left. 2 \gamma_- \lambda_1 + 2 \gamma_+ \lambda_1 + 2 \gamma_- \lambda_2 + 2 \gamma_+ \lambda_2 + 2 \lambda_1 \lambda_2 + 2 \gamma_+ \lambda_3 + 2 \lambda_{-1} \lambda_3 + 2 \lambda_1 \lambda_3 + 2 \lambda_2 \lambda_3) \right) + \\
& (-1 + y) \left( - \left( \left( e^{t (-\gamma_- - \lambda_3)} \left( \frac{1}{2} \left( -\lambda_{-1} + \lambda_1 - \lambda_2 - \sqrt{(-4 \lambda_1 \lambda_2 + (\lambda_{-1} + \lambda_1 + \lambda_2)^2)} \right) \right) \right) \right. \right. \\
\end{aligned}$$



$$2\gamma_- \lambda_1 + 2\gamma_+ \lambda_1 + 2\gamma_- \lambda_2 + 2\gamma_+ \lambda_2 + 2\lambda_1 \lambda_2 + 2\gamma_+ \lambda_3 + 2\lambda_- \lambda_3 + 2\lambda_1 \lambda_3 + 2\lambda_2 \lambda_3) \Bigg) \Bigg\} \Bigg)$$

## Simpler examples, more details

```

In[1]:= MomentEquations[{"X" → 0}, {k}, & /@ Range[4] // Column
Ex'[t] == -k Ex[t]
Ex2'[t] == -k Ex[t] - 2 k (-Ex[t] + Ex2[t])
Out[1]= Ex3'[t] == -k Ex[t] - 6 k (-Ex[t] + Ex2[t]) - 3 k (2 Ex[t] - 3 Ex2[t] + Ex3[t])
Ex4'[t] == -k Ex[t] - 14 k (-Ex[t] + Ex2[t]) -
18 k (2 Ex[t] - 3 Ex2[t] + Ex3[t]) - 4 k (-6 Ex[t] + 11 Ex2[t] - 6 Ex3[t] + Ex4[t])

In[2]:= MomentEquations[reac = {2 "Y" ← "X" + "Y" → 2 "X"}, {1, 2}, {k1, k2}]
Out[2]= ExY2'[t] == 3 k1 (Ex2Y[t] - ExY[t]) - k2 (Ex2Y[t] - ExY[t]) - k1 (ExY2[t] - ExY[t]) -
k2 (ExY2[t] - ExY[t]) - k1 ExY[t] - k2 ExY[t] + 2 k1 (Ex2Y2[t] - Ex2Y[t] - ExY2[t] + ExY[t]) -
2 k2 (Ex2Y2[t] - Ex2Y[t] - ExY2[t] + ExY[t]) + k2 (-3 ExY2[t] + ExY3[t] + 2 ExY[t]) -
k1 (-3 ExY2[t] + ExY3[t] + 2 (ExY2[t] - ExY[t]) + 2 ExY[t])

In[3]:= MasterEquation[reac, {k2, k1}, {t, x, y}, P]
Out[3]= {P^(1,0,0)[t, x, y] == (-1+x) (1+y) P[t, -1+x, 1+y] k1 +
(1+x) (-1+y) P[t, 1+x, -1+y] k2 - P[t, x, y] (x y k1 + x y k2)}

In[4]:= pgfe = ProbabilityGeneratingFunctionEquation[reac, {k2, k1}, {t, z1, z2}, G]
Out[4]= G^(1,0,0)[t, z1, z2] == k1 (z1^2 - z1 z2) G^(0,1,1)[t, z1, z2] + k2 (-z1 z2 + z2^2) G^(0,1,1)[t, z1, z2]

In[5]:= StationaryProbabilityDistributionEquation[reac, {k2, k1}, {x, y}, P]
Out[5]= {(-1+x) (1+y) P[-1+x, 1+y] k1 +
(1+x) (-1+y) P[1+x, -1+y] k2 - P[x, y] (x y k1 + x y k2) == 0}

In[6]:= RSolve[
StationaryProbabilityDistributionEquation[reac, {k2, k1}, {x, y}, P], P[x, y], {x, y}]
Out[6]= {P[x, y] → -C[1] [x + y] x y + ((k1 x - k2 x) k2^2 C[2] [x + y]) / (x y k1 (k1 - k2))}

In[7]:= dm = DeterministicModel[reac, {k2, k1}, {x, y}, t][[1]]
Out[7]= {x'[t] == k1 x[t] y[t] - k2 x[t] y[t], y'[t] == -k1 x[t] y[t] + k2 x[t] y[t]}

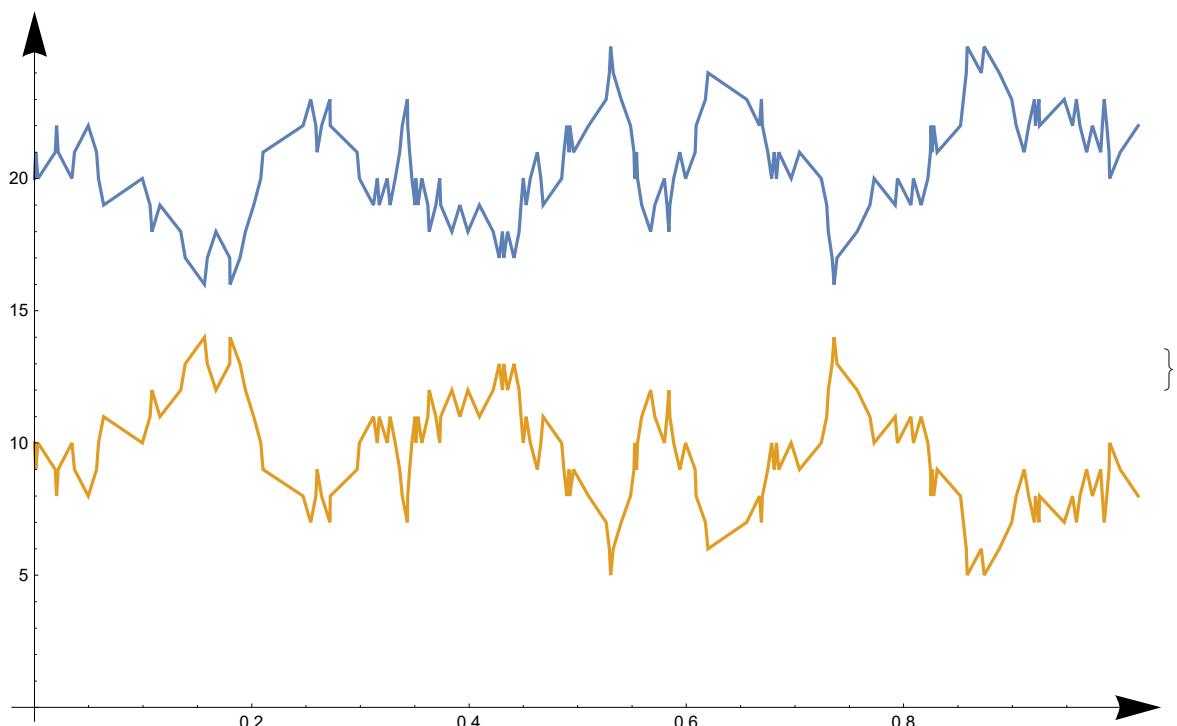
In[8]:= dm[[1]] /. {y[t] → x0 + y0 - x[t]} // Simplify
Out[8]= x'[t] == (k1 - k2) (x0 + y0 - x[t]) x[t]

```

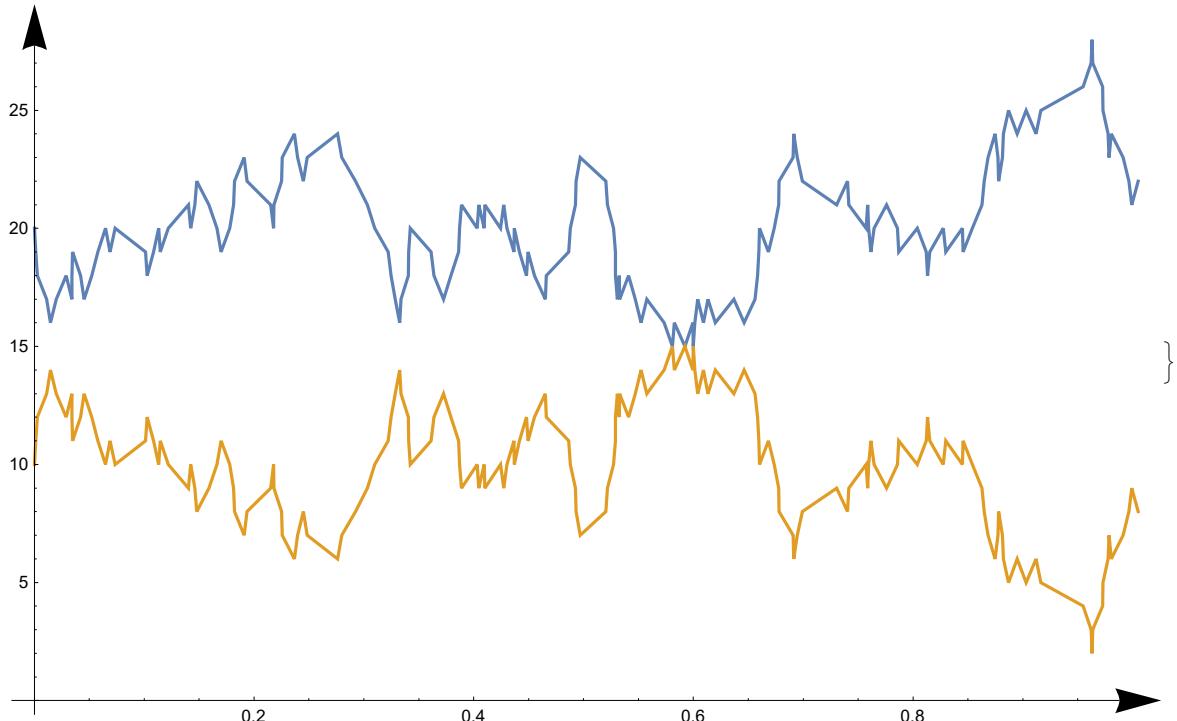
## ■ Simulation

```
SeedRandom[500];
sp1 = SimulationPlot[{"X" ≠ "Y"}, {3.8, 8.3},
{20, 10}, 1, Verbose → True, ImageSize → 600] // Timing
```

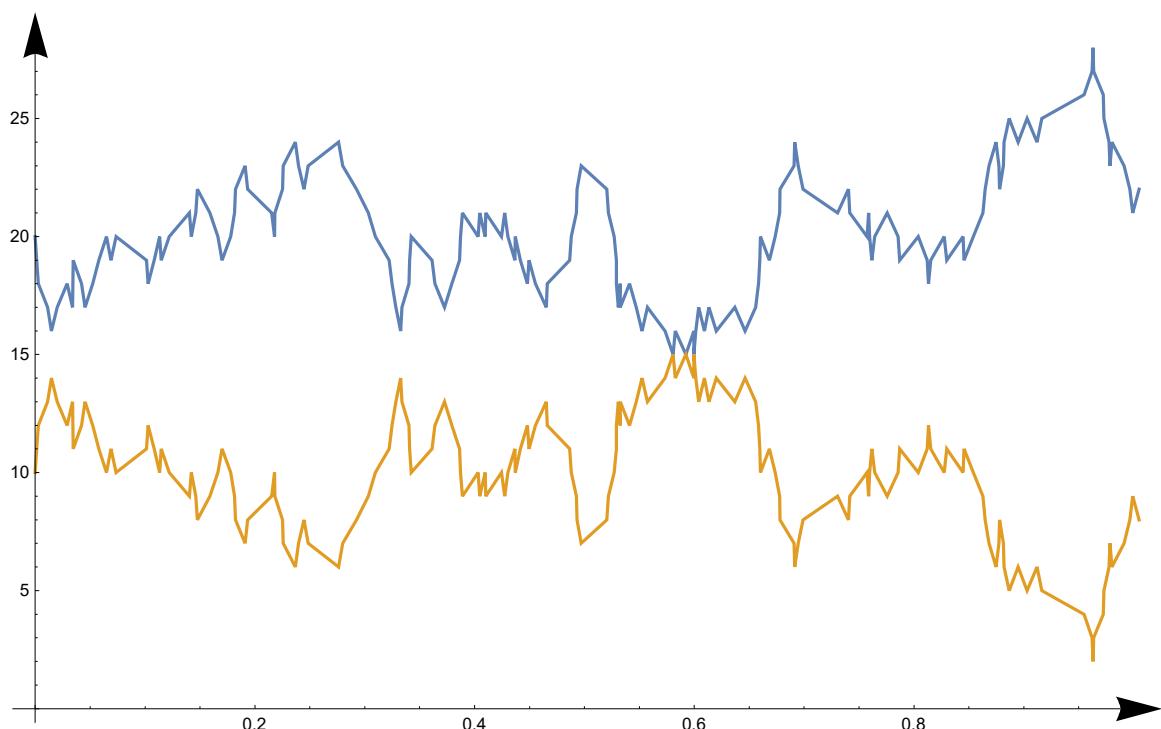
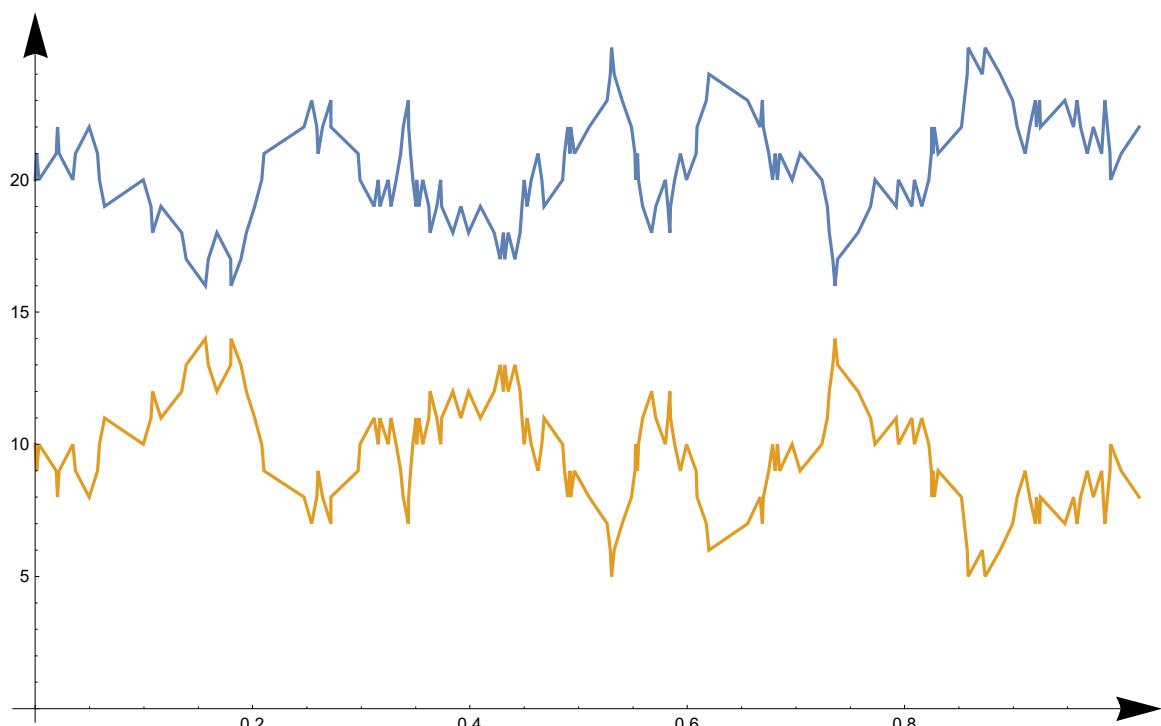
```
{0.140625,
```



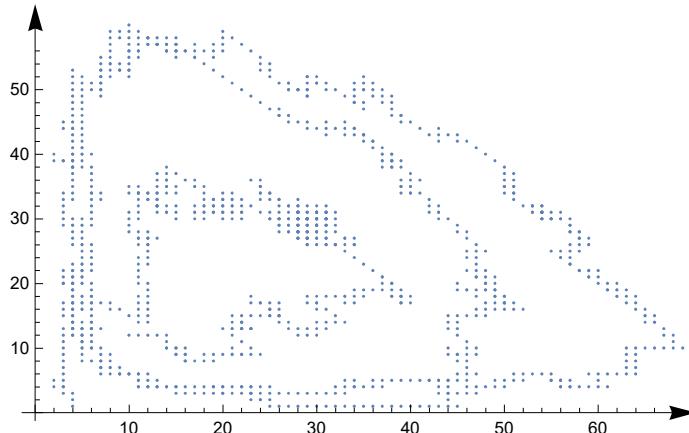
```
SeedRandom[1];
sp2 = SimulationPlot[{"X" ≠ "Y"}, {3.8, 8.3},
{20, 10}, 1, Verbose → True, ImageSize → 600] // Timing
{0.09375,
```



```
differentsimul = Column[Last /@ {sp1, sp2}]
```



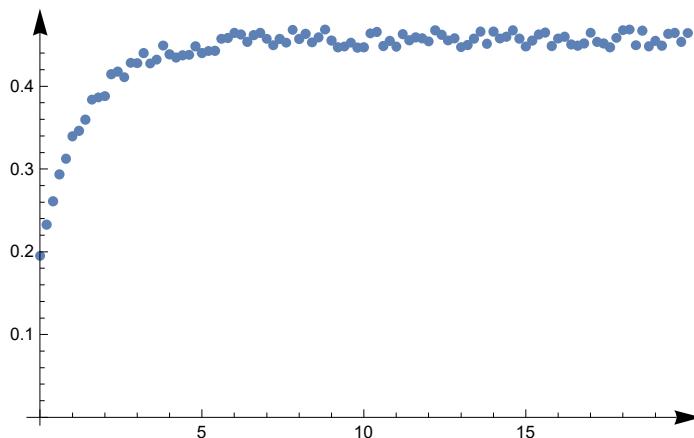
```
SeedRandom[1];
SimulationPlot2D[{"Lotka-Volterra"}, {2, 0.1, 2}, {10, 12}, 10, ExternalSpecies → {"A", "B"}]
```



## Inverse problems

### ■ Parameter estimation

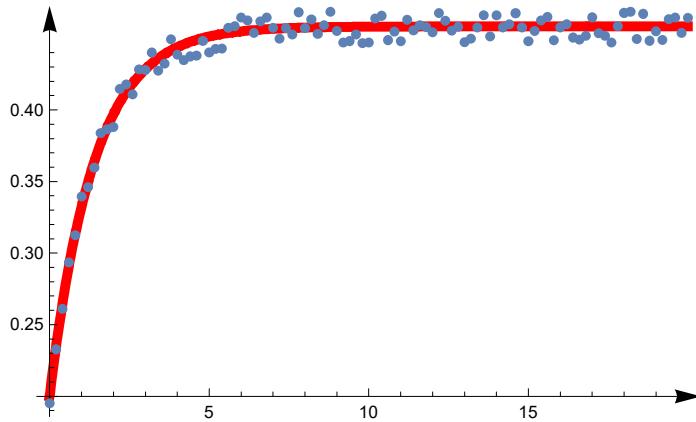
```
ClearAll[sol, times, data];
sol := First[ReplaceAll @@ Concentrations[{0 ↔ "X"}, {0.33, 0.72}, {0.2}, {0, 20}]];
times = N[Range[0, 100] / 5]; SeedRandom[100];
data = Transpose[{times, (sol /. t → times) (1 + RandomReal[{-0.025, 0.025}, 101])}];
simdata = ListPlot[data, PlotRange → All, PlotStyle → Directive[PointSize[0.015]]]
```



```

ClearAll[a, b, model2, nlm2, pl2];
model2[a_?NumberQ, b_?NumberQ, c_?NumberQ] :=
  (model1[a, b, c] = Function[t, Evaluate[First[
    ReplaceAll@@Concentrations[{0 \[leftrightarrow] "X"}, {0.33, 0.72}, {0.2}, {0, 20}]]]]));
nlm2 = NonlinearModelFit[data, model2[a, b, c][t], {{a, .1}, {b, .1}, {c, 1}}, t]; //
  Quiet
pl2 = Plot[Evaluate[nlm2[t]], {t, 0, 20}, PlotRange \[Rule] All,
  PlotStyle \[Rule] Directive[Red, Thickness[0.015]], Epilog \[Rule] First@simdata,
  ImagePadding \[Rule] {{Automatic, Automatic}, {10, 10}}]

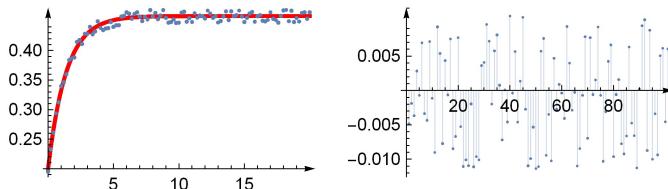
```



```

lp2 = ListPlot[nlm2["FitResiduals"], Filling \[Rule] Axis,
  ImagePadding \[Rule] {{Automatic, Automatic}, {10, 10}}];
GraphicsRow[{pl2, lp2}]

```



```

nlm2["Properties"]
{AdjustedRSquared, AIC, AICc, ANOVATable, ANOVATableDegreesOfFreedom,
 ANOVATableEntries, ANOVATableMeanSquares, ANOVATableSumsOfSquares,
 BestFit, BestFitParameters, BIC, CorrelationMatrix, CovarianceMatrix,
 CurvatureConfidenceRegion, Data, EstimatedVariance, FitCurvatureTable,
 FitCurvatureTableEntries, FitResiduals, Function, HatDiagonal, MaxIntrinsicCurvature,
 MaxParameterEffectsCurvature, MeanPredictionBands, MeanPredictionConfidenceIntervals,
 MeanPredictionConfidenceIntervalTable, MeanPredictionConfidenceIntervalTableEntries,
 MeanPredictionErrors, ParameterBias, ParameterConfidenceIntervals,
 ParameterConfidenceIntervalTable, ParameterConfidenceIntervalTableEntries,
 ParameterConfidenceRegion, ParameterErrors, ParameterPValues, ParameterTable,
 ParameterTableEntries, ParameterTStatistics, PredictedResponse, Properties,
 Response, RSquared, SingleDeletionVariances, SinglePredictionBands,
 SinglePredictionConfidenceIntervals, SinglePredictionConfidenceIntervalTable,
 SinglePredictionConfidenceIntervalTableEntries,
 SinglePredictionErrors, StandardizedResiduals, StudentizedResiduals}

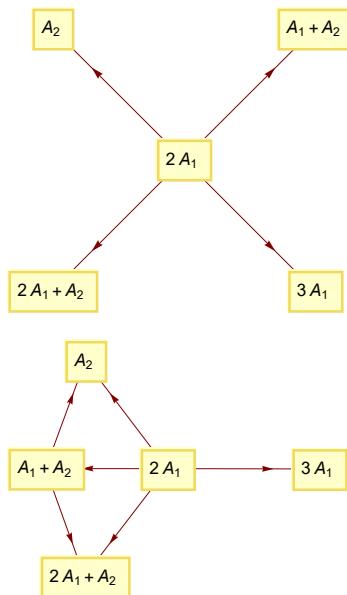
```

## ■ Finding reaction to a given (kinetic) differential equation

### Reactions with the same deterministic model

#### Different structures

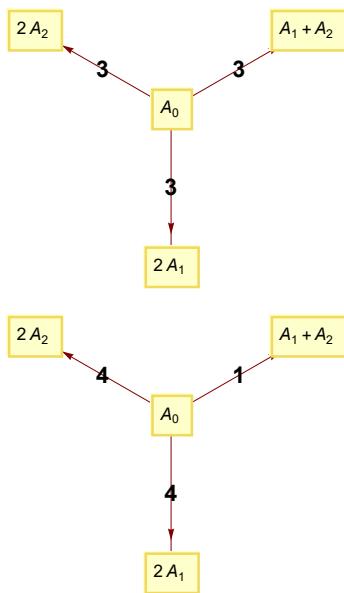
```
Column[ShowFHJGraph[##] & /@ {{2 A1 → A2, 2 A1 → 2 A1 + A2, 2 A1 → 3 A1, 2 A1 → A1 + A2}, {2 A1 → A2, 2 A1 → 2 A1 + A2, 2 A1 → 3 A1, 2 A1 → A1 + A2, A1 + A2 → 2 A1 + A2, A1 + A2 → A2} }]
```



```
MapThread[RightHandSide[#, #2, {x, y}] &,
{{{{2 A1 → A2, 2 A1 → 2 A1 + A2, 2 A1 → 3 A1, 2 A1 → A1 + A2}, {2 A1 → A2, 2 A1 → 2 A1 + A2, 2 A1 → 3 A1, 2 A1 → A1 + A2, A1 + A2 → 2 A1 + A2, A1 + A2 → A2} }, {Array[1 &, 4], {1, 19/10, 1/10, 1/10, 1/10, 1/10}}}]
{{{-2 x2, 3 x2}, {-2 x2, 3 x2}}}
```

## Different reaction rate coefficients

```
Column[{ShowFHJGraph[craciunpantea = {"A0" → 2 "A2", "A0" → 2 "A1", "A0" → "A1" + "A2"}],  
Style[#, Bold, 12] & /@ {3, 3, 3}],  
ShowFHJGraph[craciunpantea, Style[#, Bold, 12] & /@ {4, 4, 1}]}]
```



```
RightHandSide[{craciunpantea}, #] & /@ {{3, 3, 3}, {4, 4, 1}}  
{ {-9 cA0, 9 cA0, 9 cA0}, {-9 cA0, 9 cA0, 9 cA0} }
```

## Reaction with no effect on the rhs

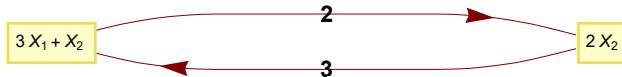
```
DeterministicModel[{2 "Y" ← "X" + "Y" → 2 "X"}, {k, k}, {x, y}][[1]]  
{x'[t] == 0, y'[t] == 0}
```

Inducing reactions with best properties (zero deficiency, weak reversibility etc.)

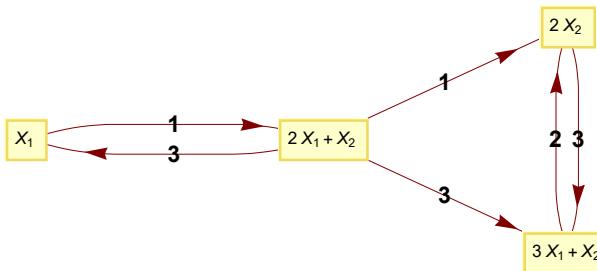
If you want to find them, start here:

```
Hyperlink["Szederkényi",  
 "https://scholar.google.hu/citations?user=-f46rMcAAAAJ&hl=hu&oi=ao"]  
Szederkényi
```

```
dyneqsteps1 = {"X1"  $\rightleftharpoons$  2 "X1" + "X2", 2 "X2"  $\rightleftharpoons$  3 "X1" + "X2"};
dyneq1 = ShowFHJGraph[dyneqsteps1, Style[#, Bold, 12] & /@ {1, 2, 3, 2}]
```



```
dyneqsteps2 = {"X1"  $\rightleftharpoons$  2 "X1" + "X2"  $\rightarrow$  2 "X2"  $\rightleftharpoons$  3 "X1" + "X2"  $\leftarrow$  2 "X1" + "X2"};
dyneq2 = ShowFHJGraph[dyneqsteps2, Style[#, Bold, 12] & /@ {1, 3, 1, 3, 2, 3}]
```



```
szeder1 = {2 A1  $\rightarrow$  A2, 2 A1  $\rightarrow$  A1 + A2, 2 A1  $\rightarrow$  3 A1, 2 A1  $\rightarrow$  2 A1 + A2} ;
s1 = ShowFHJGraph[szeder1, Array[1 &, 4], DirectedEdges -> True, VertexLabeling -> True]
figureexp["TothNagyPapp-Fig11.5A", s1]

szeder2 =
{2 A1  $\rightarrow$  A2, 2 A1  $\rightarrow$  A1 + A2, 2 A1  $\rightarrow$  3 A1, 2 A1  $\rightarrow$  2 A1 + A2, A1 + A2  $\rightarrow$  A2, A1 + A2  $\rightarrow$  2 A1 + A2} ;
s2 = ShowFHJGraph[szeder2, {1, 0.1, 0.1, 1.9, 0.1, 0.1},
DirectedEdges -> True, VertexLabeling -> True]

origr = {X + Y  $\rightleftharpoons$  X + 2 Y  $\rightleftharpoons$  2 X + 3 Y  $\rightarrow$  2 Y  $\leftarrow$  X + Y, 2 Y  $\rightleftharpoons$  X + 2 Y} ;
orig = ShowFHJGraph[origr,
Style[#, Bold, 16] & /@ ({"k1", "k2", "k5", "k4", "k8", "k3", "k7", "k6"} /.
{"k1"  $\rightarrow$  1, "k2"  $\rightarrow$  "11/10", "k5"  $\rightarrow$  "11/10", "k4"  $\rightarrow$  1, "k8"  $\rightarrow$  1,
"k3"  $\rightarrow$  1, "k7"  $\rightarrow$  3, "k6"  $\rightarrow$  "1/10"}, ImageSize  $\rightarrow$  1300]

sparser = {X + Y  $\rightleftharpoons$  X + 2 Y  $\rightleftharpoons$  2 X + 3 Y  $\rightarrow$  2 Y  $\leftarrow$  X + Y, 2 Y  $\rightarrow$  X + 2 Y} ;
sparse = ShowFHJGraph[sparser,
Style[#, Bold, 16] & /@ ({"k1", "k2", "k3", "k4", "k5", "k6", "k7"} /.
{k1"  $\rightarrow$  1, "k2"  $\rightarrow$  1, "k3"  $\rightarrow$  1, "k4"  $\rightarrow$  1, "k5"  $\rightarrow$  1, "k6"  $\rightarrow$  1, "k7"  $\rightarrow$  3}), ImageSize  $\rightarrow$  1000]

denser = {2 X + 3 Y  $\rightleftharpoons$  X + Y  $\rightleftharpoons$  X + 2 Y  $\rightleftharpoons$  2 X + 3 Y  $\rightleftharpoons$  2 Y  $\rightleftharpoons$  X + Y, 2 Y  $\rightleftharpoons$  X + 2 Y} ;
dense = ShowFHJGraph[denser, Style[#, Bold, 16] & /@
({{"k1", "k2", "k3", "k4", "k5", "k6", "k7", "k8", "k9", "k10", "k11", "k12"} /.
{"k1"  $\rightarrow$  "3/10", "k2"  $\rightarrow$  "3/10", "k3"  $\rightarrow$  "1/10", "k4"  $\rightarrow$  "11/10", "k5"  $\rightarrow$  "11/10",
"k6"  $\rightarrow$  "1/10", "k7"  $\rightarrow$  "13/10", "k8"  $\rightarrow$  "29/30", "k9"  $\rightarrow$  "29/30",
"k10"  $\rightarrow$  "13/10", "k11"  $\rightarrow$  "1/10", "k12"  $\rightarrow$  "1/10"}, ImageSize  $\rightarrow$  800]
```

## Identifiability and Uniqueness Questions

No compartmental system induces this kinetic differential equation

```
RightHandSide[{"Y" → 3 "X", "X" → 3 "Y"}, {1, 1}, {x, y}]
{-x + 3 y, 3 x - y}
```

# Programs

## ■ Other programs

[Hyperlink\["Demonstrations", "http://demonstrations.wolfram.com/"\]](http://demonstrations.wolfram.com/)

Ferreira MMC, Ferreira WCJ, Lino ACS, Porto MEG: Uncovering oscillations, complexity, and chaos in chemical kinetics using Mathematica. *J Chem Educ* **76**(6) (1999), 861–866.

FrancI MM (2000) Introduction to the use of numerical methods in chemical kinetics.

[Hyperlink\["FrancI", "http://ckw.phys.ncku.edu.tw/public/pub/WebSources/MathSource/library.wolfram.com/infocenter/MathSource/791/index.html"\]](http://ckw.phys.ncku.edu.tw/public/pub/WebSources/MathSource/library.wolfram.com/infocenter/MathSource/791/index.html)

FrancI

Hollis S (2004) Reaction-diffusionlab.m.

[Hyperlink\["Hollis", "http://www.math.armstrong.edu/faculty/hollis/mmade/RDL"\]](http://www.math.armstrong.edu/faculty/hollis/mmade/RDL)

Kyurkchiev N, Markov S, Mincheva M.: Analysis of biochemical mechanisms using Mathematica with applications. *Serdica J. Computing* 10 (2016), No 1, 63–78

Mulquiney PM, Kuchel PW (2003) Modelling metabolism with Mathematica. CRC Press, Boca Raton, FL, URL

[Hyperlink\["Kuchel, includes CD-ROM",
 "http://www.bioeng.auckland.ac.nz/MCF/invited\\_users/philip\\_kuchel.htm"\]](http://www.bioeng.auckland.ac.nz/MCF/invited_users/philip_kuchel.htm)

Sánchez G (2005) Biokmod: A Mathematica toolbox for modeling biokinetic systems. *Mathematica in Education and Research* 10(2):1–34, URL

[Hyperlink\["Sanchez", "http://diarium.usal.es/guillermo/biokmod/"\]](http://diarium.usal.es/guillermo/biokmod/)

## ■ Information about our ReactionKinetics.wl

[Hyperlink\["Springer", "http://extras.springer.com"\]](http://extras.springer.com)

Springer

```
In[8]:= OpenReactionKineticsPalette[]

Out[8]= NotebookObject[ ReactionKinetics Palette]
```

```
In[9]:= Names["ReactionKinetics`*"]

Out[9]= {AbsoluteConcentrationRobustness, AcyclicVolpertGraphQ, AtomConservingQ, Atoms,
AtomsQ, AvogadrosNumber, BioModels, BioReactions, Bipartite, CHEMKINExport,
CHEMKINImport, CombinatorialMoments, ComplexColors, ComponentwiseLessEqualQ,
Concentrations, Conditions, ContejeanDevie, CoveringDecompositionSet,
Decompositions, DeleteAutocatalysis, DependencyGraph, DetailedBalanced,
DeterministicModel, EdgeLabeled, EigensystemJacobian, ElementaryReactions,
ExternalSpecies, FastSelection, Filter, FilterReactions, FixedStepSize,
FormattedOutput, FromAtomMatrix, FromStoichiometry, GammaLeftNullSpace,
GeneratedRateCoefficient, GetReaction, GreedySelection, Highlight, LPBased,
MassAction, MassActionKinetics, MassConservationRelations, MasterEquation,
MaxFHJStronglyConnectedComponents, MaxFHJWeaklyConnectedComponents, MaxIteration,
Memo, MinFHJStronglyConnectedComponents, MinFHJWeaklyConnectedComponents,
MinimalDecompositions, Models, MolarGasConstant, MomentEquations, Numbered,
ObjectiveFunction, Obligatory, Omittable, OpenReactionKineticsNamesPalette,
OpenReactionKineticsPalette, OriginalSelection, PlotFunction, Positivity,
Preprocess, ProbabilityGeneratingFunctionEquation, ReactionRatesNotebook,
Reactions, ReactionsData, ReversibleFHJRepresentation, ReversibleQ, RightHandSide,
SelectMinimalDecompositions, ShowDependencyGraph, ShowFHJGraph, ShowVolpertGraph,
Side, Simulation, SimulationPlot, SimulationPlot2D, SimulationPlot3D,
SolveProbabilityGeneratingFunctionEquation, Species, StationaryPoints,
StationaryProbabilityDistributionEquation, StronglyConnectedComponentsColors,
SubgraphHighlight, TimeLimit, ToAtomMatrix, ToCanonicalForm, ToReversible,
VolpertIndexing, WeaklyReversibleQ, ZeroVectorQ, $ReactionKinetics,
$ReactionKineticsPackageLoaded, $ReactionKineticsVersionNumber}
```

```
In[10]:= Information["ReactionKinetics`*"]
```

▼ ReactionKinetics`

AbsoluteConcentrationRobustness

AcyclicVolpertGraphQ

AtomConservingQ

Atoms

AtomsQ

AvogadrosNumber

BioModels

BioReactions

Bipartite

CHEMKINExport

CHEMKINImport

CombinatorialMoments

ComplexColors

ComponentwiseLessEqualQ

Concentrations

Conditions

ContejeanDevie

CoveringDecompositionSet

Decompositions
DeleteAutocatalysis
DependencyGraph
DetailedBalanced
DeterministicModel
EdgeLabeled
EigensystemJacobian
ElementaryReactions
ExternalSpecies
FastSelection
Filter
FilterReactions
FixedStepSize
FormattedOutput
FromAtomMatrix
FromStoichiometry
GammaLeftNullSpace
GeneratedRateCoefficient
GetReaction
GreedySelection
Highlight
LPBased
MassAction
MassActionKinetics
MassConservationRelations
MasterEquation
MaxFHJStronglyConnectedComponents
MaxFHJWeaklyConnectedComponents
MaxIteration
Memo
MinFHJStronglyConnectedComponents
MinFHJWeaklyConnectedComponents
MinimalDecompositions
Models
MolarGasConstant
MomentEquations
Numbered
ObjectiveFunction
Obligatory
Omittable
OpenReactionKineticsNamesPalette
OpenReactionKineticsPalette
OpenReactionKineticsNames

```

OriginalSelection
PlotFunction
Positivity
Preprocess
ProbabilityGeneratingFunctionEquation
ReactionRatesNotebook
Reactions
ReactionsData
ReversibleFHJRepresentation
ReversibleQ
RightHandSide
SelectMinimalDecompositions
ShowDependencyGraph
ShowFHJGraph
ShowVolpertGraph
Side
Simulation
SimulationPlot
SimulationPlot2D
SimulationPlot3D
SolveProbabilityGeneratingFunctionEquation
Species
StationaryPoints
StationaryProbabilityDistributionEquation
StronglyConnectedComponentsColors
SubgraphHighlight
TimeLimit
ToAtomMatrix
ToCanonicalForm
ToReversible
VolpertIndexing
WeaklyReversibleQ
ZeroVectorQ
$ReactionKinetics
$ReactionKineticsPackageLoaded
$ReactionKineticsVersionNumber

```

---

**AbsoluteConcentrationRobustness[{reactions},options]**  
checks the Shinar–Feinberg conditions for the reaction.

Attributes[AbsoluteConcentrationRobustness] = {Protected, ReadProtected}

Options[AbsoluteConcentrationRobustness] := {TimeLimit → 120, ExternalSpecies → {}}

In[11]:= **? Concentrations**

Concentrations[{reactions},rates,initvalues,vars,options] attempts to solve the induced kinetic differential equation endowed with some kinetics described by the rates. Initial concentrations are given by initvalues, vars contain the names of the independent variables. This way Concentrations uses DSolve, accepts symbolic parameters, and tries to return the symbolic solution to the equation (i.e. concentrations versus time curves).  
 Concentrations[{reactions},rates,initvalues,{t0,t1},vars,options]  
 uses NDSolve to give the numerical solution from t0 to t1.  
 Concentrations[{reactions},arrheniuscoeffs,enthalpies,c0,V,Ta,tres,p1,p2,initvalues,{t0,t1},options]  
 tries to solve the general evolution equation for reactions involving temperature effects. In this case the kinetics is assumed to be of the mass action type, the temperature dependence is allowed to be of the generalized Arrhenius type. The Arrhenius triples are given by arrheniuscoeffs.

In[12]:= **? Simulation**

Simulation[{reactions},rratecoeffs,init,maxtime,options] simulates the induced kinetic Markov process of the reaction endowed with stochastic mass action type kinetics with stochastic reaction rate coefficients given by rratecoeffs. The initial state are given by init, while the stopping time is specified by maxtime. Options include ExternalSpecies, FixedStepSize, Tolerance (for tau-leaping), MaxIteration, Method, Verbose and Volume. The following methods can be used: "Direct" (default), "FirstReaction", "NextReaction", "ExplicitTau-Leaping", "ImplicitTau-Leaping" and "TrapezoidalTau-Leaping".

**? SimulationPlot**

SimulationPlot[{reactions},rratecoeffs,init,maxtime,options] simulates and plots the trajectories of species, where the induced kinetic Markov endowed with stochastic mass action type kinetics with stochastic reaction rate coefficients given by rratecoeffs. Options include ExternalSpecies, PlotFunction (default is "ListLinePlot") and Species (default is All).

In[13]:= **Options [ShowFHJGraph]**

Out[13]= {ExternalSpecies → {}, ComplexColors → {}, PlotFunction → GraphPlot,  
 StronglyConnectedComponentsColors → {}, Numbered → False}

**? ShowFHJGraph**

In[14]:= **Models**

Out[14]= {Autocatalator, Belousov-Zhabotinsky, Briggs-Rauscher, Brusselator, Chapman cycle, Clarke, Colquhoun, Consecutive, Decomposition, DeYoung-Keizer, Dimerization, Edelstein, Eigen, Érdi-Ropolyi, Explodator, Horn-Jackson, Feinberg-Horn I, Feinberg-Horn II, FKN mechanism, Frank, Goldbeter, Hudson-Rössler, Huxel, Hyver, Inflow, Ivanova, Kliemann, Leonard-Reichl, Lotka, Lotka-Volterra, Michaelis-Menten, Mole, Ogg, Oregonator, Outflow, Petri, Robertson, Schlögl, Simple birth-death, Triangle, Turányi-Györgyi-Field, Vaiman, Verhulst, Volpert, Wegscheider, Willamowski-Rössler}

In[15]:= **BioModels**

Out[15]= {Calvin Cycle (Photosynthesis), Glycolysis, Glyoxylate Cycle, SzentGyörgyi-Krebs Cycle}

In[16]:= **Reactions**

Out[16]=  $\left\{ \begin{array}{l} \text{Autocatalator} \rightarrow \{A \rightarrow X, X \rightarrow Y, X + 2Y \rightarrow 3Y, Y \rightarrow P\}, \\ \text{Belousov-Zhabotinsky} \rightarrow \left\{ H + X + Y \rightarrow 2V, A + 2H + Y \rightarrow V + X, 2X \rightarrow V, \right. \\ \left. A + H + \frac{X}{2} \rightarrow X + Z, X + Z \rightarrow \frac{X}{2}, M + Z \rightarrow Q, V + Z \rightarrow Y, V \rightarrow Y, X \rightarrow 0, Y \rightarrow 0, Z \rightarrow 0, V \rightarrow 0 \right\}, \\ \text{Briggs-Rauscher} \rightarrow \{CH_2(CO_2H)2 + H \rightarrow 2H_2O + ICH(CO_2H)2 + 2O_2, \\ H \rightarrow 2H_2O + IO_3^- \rightarrow 2H_2O + HIO + 2O_2, CH_2(CO_2H)2 + HIO \rightarrow H_2O + ICH(CO_2H)2, \\ 2H \rightarrow IO_3^- \rightarrow HIO + HIO_2, H \rightarrow HIO_2 + I^- \rightarrow 2HIO, \\ H_2O_2 + HIO \rightarrow H \rightarrow H_2O + I^- + O_2, H \rightarrow HIO_2 + IO_3^- \rightarrow H_2O + 2IO_2, \\ H_2O + IO_2 + Mn^{2+} \rightarrow HIO_2 + Mn(OH)_2^-, H_2O_2 + Mn^{2+} \rightarrow H_2O + H_2O + Mn^{2+}, \\ 2H_2O \rightarrow H_2O_2 + O_2, 2HIO_2 \rightarrow H \rightarrow HIO + IO_3^-, \\ H \rightarrow HIO + I^- \rightarrow H_2O + I_2, CH_2(CO_2H)2 + I_2 \rightarrow H \rightarrow I^- + ICH(CO_2H)2\}, \\ \text{Brusselator} \rightarrow \{X \leftarrow A, B + X \rightarrow D + Y, 2X + Y \rightarrow 3X, X \rightarrow E\}, \\ \text{Chapman cycle} \rightarrow \{O_2 \rightarrow 2O, M + O \rightarrow O_2, O_3 \rightarrow O + O_2, O + O_3 \rightarrow 2O_2\}, \\ \text{Clarke} \rightarrow \{A + T \rightarrow M + S, H + T + Y \rightarrow S + V, H + X + Y \rightarrow 2T, B + 2H + Y \rightarrow T + X, 2X \rightarrow B + H + T, \\ B + H + X \rightarrow S + 2U, H + U + W \rightarrow X + Z, S + U + Z \rightarrow B + 2H + W, A + V \rightarrow H + M + Y, \\ A + 2S + 6Z \rightarrow 6H + P + 2Q + 6W, M + 2S + 4Z \rightarrow 5H + P + 2Q + 4W + Y, P + T \rightarrow H + Q + S + Y\}, \\ \text{Colquhoun} \rightarrow \{AR \rightleftharpoons A + R \rightleftharpoons RA \rightleftharpoons A + RA\}, \text{Consecutive} \rightarrow \{A \rightarrow B \rightarrow C\}, \\ \text{Decomposition} \rightarrow \{C \rightarrow A + B\}, \\ \text{DeYoung-Keizer} \rightarrow \{S(000) \rightleftharpoons S(001) \rightleftharpoons S(100) \rightleftharpoons S(010), \\ S(100) \rightleftharpoons S(101) \rightleftharpoons S(110), S(001) \rightleftharpoons S(101) \rightleftharpoons S(011), \\ S(011) \rightleftharpoons S(010) \rightleftharpoons S(111), S(110) \rightleftharpoons S(010) \rightleftharpoons S(111), S(111) \rightleftharpoons S(101)\}, \\ \text{Dimerization} \rightarrow \{A + B \rightarrow C\}, \text{Edelstein} \rightarrow \{X \rightleftharpoons 2X, X + Y \rightleftharpoons Z \rightleftharpoons Y\}, \\ \text{Eigen} \rightarrow \{X \rightarrow 0, X \rightarrow 2X\}, \text{Érdi-Ropolyi} \rightarrow \{R + 4T \rightleftharpoons T4R1 \rightleftharpoons T4R2 \rightleftharpoons T4R3\}, \\ \text{Explodator} \rightarrow \{A + X \rightarrow X (1 + a\text{Explodator}), X + Y \rightarrow Z, Z \rightarrow Y (1 + b\text{Explodator}), Y \rightarrow P\}, \\ \text{Horn-Jackson} \rightarrow \{3X \rightarrow X + 2Y \rightarrow 3Y \rightarrow 2X + Y \rightarrow 3X\}, \\ \text{Feinberg-Horn I} \rightarrow \{2J \rightarrow G, 2J \leftrightarrow H, G \rightarrow H, A + B \rightarrow G, A + B \leftrightarrow C, C \rightarrow D + E, D + E \leftrightarrow F\}, \\ \text{Feinberg-Horn II} \rightarrow \{X \rightarrow Y, X + Z \rightarrow T \rightarrow U + Y \rightarrow X + Z\}, \text{FKN mechanism} \rightarrow \\ \{Br^- + HBrO_2 \rightarrow 2HBr, Br^- + BrO_3^- \rightarrow HBrO_2 + HOBr, 2HBrO_2 \rightarrow BrO_3^- + HOBr, \\ Br^- + HOBr \rightarrow Br_2 + H_2O, BrO_3^- + HBrO_2 \rightarrow 2BrO_2 + H_2O, H_2O + HBrO_2 \rightarrow BrO_3^-, \\ Br_2 + MA \rightarrow Br^- + BrMA, 2H_2O + MA \rightarrow 2CO_2 + C, BrMA + 2H_2O \rightarrow Br^- + 2CO_2 + HCO_2H\}, \\ \text{Frank} \rightarrow \{A \rightarrow R, A \rightarrow S, A + R \rightarrow 2R, A + S \rightarrow 2S\}, \text{Goldbeter} \rightarrow \{M \rightleftharpoons 0 \rightleftharpoons C, 0 \rightleftharpoons X, C \rightarrow 0\}, \\ \text{Hudson-Rössler} \rightarrow \{P \rightarrow A, Q \rightarrow B, A + B \rightarrow 2B, B \rightarrow R, A \leftrightarrow C\}, \\ \text{Huxel} \rightarrow \{X \rightleftharpoons 2X, X + Y \rightarrow 2Y, Y \rightarrow 0\}, \text{Hyver} \rightarrow \{K1 \rightarrow A1 \rightarrow A2 \rightarrow A3 \rightarrow A4 \rightarrow A5 \rightarrow A6 \rightarrow A7, \\ A7 + B \rightarrow C + E, K \rightarrow B \rightarrow 0, A1 + E \rightarrow 0, A1 + C \rightarrow 0, \mu \rightarrow D, C + D \rightarrow 0, D + E \rightarrow 0\}, \\ \text{Inflow} \rightarrow \{0 \rightarrow X\}, \text{Ivanova} \rightarrow \{X + Y \rightarrow 2Y, Y + Z \rightarrow 2Z, X + Z \rightarrow 2X\}, \\ \text{Kliemann} \rightarrow \{X \rightleftharpoons Y, 2X \rightleftharpoons X + Y \rightleftharpoons 2Y\}, \text{Leonard-Reichl} \rightarrow \{2X \rightleftharpoons Y\}, \\ \text{Lotka} \rightarrow \{A \rightarrow X, X + Y \rightarrow 2Y, Y \rightarrow P\}, \text{Lotka-Volterra} \rightarrow \{A + X \rightarrow 2X, X + Y \rightarrow 2Y, B \leftarrow Y\}, \\ \text{Michaelis-Menten} \rightarrow \{E + S \rightleftharpoons C \rightarrow E + P\}, \text{Mole} \rightarrow \{X + Y \rightarrow 2X + 2Y, X \leftrightarrow 0 \leftrightarrow Y\}, \\ \text{Ogg} \rightarrow \{N_2O_5 \rightleftharpoons NO_2 + NO_3, NO_2 + NO_3 \rightarrow NO + NO_2 + O_2, NO + NO_3 \rightarrow 2NO_2\}, \\ \text{Oregonator} \rightarrow \{A + Y \rightarrow X, X + Y \rightarrow P, B + X \rightarrow 2X + Z, 2X \rightarrow Q, Z \rightarrow Y\}, \\ \text{Outflow} \rightarrow \{X \rightarrow 0\}, \text{Petri} \rightarrow \{C + O_2 \rightarrow CO_2, NaOH + CO_2 \rightarrow NaHCO_3, HCl + NaHCO_3 \rightarrow NaCl + CO_2 + H_2O\}, \\ \text{Robertson} \rightarrow \{A \rightarrow B, 2B \rightarrow B + C, B + C \rightarrow A + C\}, \\ \text{Schlögl} \rightarrow \{0 \rightleftharpoons X, 2X \rightleftharpoons 3X\}, \text{Simple birth-death} \rightarrow \{X \rightarrow 2X, X \rightarrow A\}, \\ \text{Triangle} \rightarrow \{A \rightarrow B \rightarrow C \rightarrow A\}, \text{Turányi-Györgyi-Field} \rightarrow \\ \{X + Y \rightarrow 2P, A + Y \rightarrow P + X, 2X \rightarrow A + P, A + X \rightarrow 2X + 2Z, X + Z \rightarrow A + 0.5X, M + Z \rightarrow Y - Z\}, \end{array} \right.$

Vaiman  $\rightarrow \{ 2 \text{K} + \text{O}_2 \rightarrow 2 \text{KO}, \text{H}_2\text{S} + \text{K} \rightarrow \text{KH}_2\text{S}, 2 \text{KH}_2\text{S} + \text{O}_2 \rightarrow 2 \text{H}_2\text{O} + \text{K} + \text{KS}_2, \text{H}_2\text{S} + \text{KO} \rightarrow \text{H}_2\text{O} + \text{KS}, \text{KH}_2\text{S} + \text{KO} \rightarrow \text{H}_2\text{O} + \text{K} + \text{KS}, 2 \text{KS} \rightarrow \text{K} + \text{KS}_2, \text{KS}_2 \rightarrow \text{K} + \text{S}_2 \}$ ,  
Verhulst  $\rightarrow \{ \text{X} \rightleftharpoons 2 \text{X} \}$ , Volpert  $\rightarrow \{ \text{X} + \text{Y} \rightarrow \text{T}, \text{Y} + \text{Z} \rightarrow \text{U} \}$ , Wegscheider  $\rightarrow \{ \text{A} \rightleftharpoons \text{B}, 2 \text{A} \rightleftharpoons \text{A} + \text{B} \}$ ,  
Willamowski-Rössler  $\rightarrow \{ \text{A} + \text{X} \rightleftharpoons 2 \text{X}, \text{X} + \text{Y} \rightleftharpoons 2 \text{Y}, \text{Q} + \text{Y} \rightleftharpoons \text{B}, \text{X} + \text{Z} \rightleftharpoons \text{C}, \text{P} + \text{Z} \rightleftharpoons 2 \text{Z} \}$

Mathematical Chemistry BMETE92MM09, 4 hours/week, 5 credits; 2018/19. II. Students from ELTE are also welcome.