

# Point-wise models in MFront

DE LA RECHERCHE À L'INDUSTRIE

# MFront training course

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Commissariat à l'énergie atomique et aux énergies alternatives - www.cea.fr

# Introduction



▶ Point-wise models (swelling due to solid or gaseous fission products, phase transition, chemical reactions) describes the local evolution of a set of *internal state variables*  $(y_j)_{j \in [1:n_y]}$  due to the evolution local thermodynamic environment described by a set of *external state variables*  $(z_j)_{i \in [1:n_x]}$ .



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- Point-wise models can be seen as a simpliest case of generic behaviours:
  - No thermodynamic force nor consistent tangent operator.



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- ▶ Up to a certain extent, point-wise models may also be used to describe the evolution of a structure :
  - Oxidation of Zircaloy pipes.
- ▶ Point-wise models can be seen as a simpliest case of generic behaviours :
  - **—** No thermodynamic force nor consistent tangent operator.
- ▶ Point-wise models can be used :
  - directly (in a PLEIADES application, in MTest or in MGIS) or used a build are a building block for models
  - as a building block for behaviours.



# **Evolution of internal state variables, notations**

► The *internal state variables*  $(y_j)_{j \in [1:n_v]}$  are packed in a vector  $\vec{Y}$ :

$$\vec{Y} = \begin{pmatrix} y_1 & \dots & y_{n_y} \end{pmatrix}^T$$

- Note that this notation does not make any assumption on tensorial nature of the state variable  $y_j$  that may a scalar, a symmetric tensor, an unsymmetric tensor, etc...



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- ▶ The external state variables  $(z_j)_{j \in [1:n_2]}$  packed in a vector  $\vec{Z}$ :

$$\vec{Z} = \begin{pmatrix} z_1 & \dots & z_{n_z} \end{pmatrix}^T$$



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- ▶ The external state variables  $(z_j)_{j \in [1:n_z]}$  packed in a vector  $\vec{Z}$ :

$$\vec{Z} = \begin{pmatrix} z_1 & \dots & z_{n_z} \end{pmatrix}^T$$

► The evolution of the internal state variables  $\vec{Y}$  is assumed to obey a standard first-order ordinary differential equation (ODE) given by :

$$\vec{Y} = \frac{d\vec{Y}}{d\tau} = \vec{G}(\vec{Y}(\tau), \vec{Z}(\tau))$$

where  $\tau$  denotes the time variable.



# Time discretization, notations

- ► Time is discretized in intervals named time steps.
  - t denotes the time at the beginning of the (considered) time step.
  - $-\Delta t$  is the time increment during the time step.
  - $-t + \Delta t$  corresponds to the time at the end of the step.



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  - **–**  $t + \Delta t$  corresponds to the time at the end of the step.
- Models in MFront assumes that the values of the external state variables at the beginning of the time step, denoted  $\vec{Z}\Big|_t$  and their increments during the time step, denoted  $\Delta \vec{Z}$ , are known.
  - Since only  $\vec{Z}\Big|_t$  and  $\Delta \vec{Z}$  are known, the evolution of  $\vec{Z}$  during the time step can only be described using a linear interpolation :

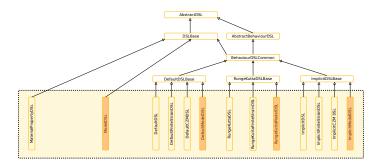
$$\left. \vec{Z} \right|_{t+\theta \, \Delta \, t} = \left. \vec{Z} \right|_{t} + \theta \, \Delta \, \vec{Z}$$

where  $\theta$  is a number in the range [0, 1]

 This assumption generally implies that high order algorithms are generally useless.



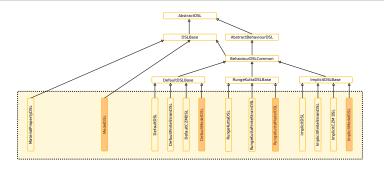
# Domain specific languages related to models



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- ▶ The Model DSL is historically the first DSL dedicated to models.
- ▶ The DefaultModel, the RungeKuttaModel and the ImplicitModel have been introduced in recent versions of MFront as a consequence of the work on generic behaviours.
  - Those DSLs are much more powerful than the Model DSL.
  - Their conventions are more consistent with the rest of MFront.
  - Alas, there is no interface for the PLEIADES architecture nor licos yet.

# A simple chemical reaction

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➤ To illustrate the implementation of models with 'MFront', let us consider a system of two chemical species A and B whose evolution is given by the following reaction:

$$A \stackrel{k_1}{\rightleftharpoons} B$$



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$$A \stackrel{k_1}{\underset{k_2}{\rightleftharpoons}} B$$

▶ The molar concentrations [A] and [B] thus satisfies :

$$\begin{cases} \frac{d[A]}{d\tau} = k_2[B](\tau) - k_1[A](\tau) \\ \frac{d[B]}{d\tau} = k_1[A](\tau) - k_2[B](\tau) \end{cases}$$



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- ▶ [A] + [B] is constant (*conservation of mass*).
- ► Hence, the evolution of the system is driven by the following condensed equation :

$$\frac{d[A]}{d\tau} = k_2([A]|_t + [B]|_t) - (k_1 + k_2)[A](\tau)$$



mentation with the Model DSL. First tests



# **Constant reaction rate coefficients**

Reaction rate coefficients	Value
k <sub>1</sub>	$\frac{1}{60} s^{-1}$
k <sub>2</sub>	$\frac{1}{120}  s^{-1}$

▶ If the reaction rate coefficients  $k_1$  and  $k_2$  are assumed constant, a closed formed solution is given by :

$$[A]|_{t+\Delta t} = \frac{B}{K} + \left(\frac{K [A]|_{t} - B}{K}\right) \exp\left(-K \Delta t\right)$$

with: 
$$B = k_2 ([A]|_t + [B]|_t)$$
 and  $K = k_1 + k_2$ .



# A first implementation

```
@DSL Model:
@Model ChemicalReaction1:
@Author Thomas Helfer:
@Date 09/07/2022:
@UseQt true;
@UnitSystem SI:
//! molar concentration of species A
@StateVariable quantity<real, 0, 0, 0, 0, 0, 1> ca:
ca.setEntryName("MolarConcentrationOfSpeciesA");
ca.setDepth(1);
//! molar concentration of species B
@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 1> cb;
cb.setEntryName("MolarConcentrationOfSpeciesB");
cb.setDepth(1);
//! rate coefficient of the reaction transforming species A to species B
k1.setEntryName("ReactionRateCoefficientAB");
//! rate coefficient of the reaction transforming species B to species A
@Parameter frequency k2 = 0.0083333333333333333;
k2.setEntryName("ReactionRateCoefficientBA"):
@Function ChemicalReaction {
 const auto B = k2 * (ca 1 + cb 1);
 const auto K = k1 + k2
 const auto e = \exp(-K * dt);
 ca = ca \ 1 * e + (B / K) * (1 - e);
 cb = ca 1 + cb 1 - ca:
```



### //! molar concentration of species A

@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 0, 1> ca; ca.setEntryName("MolarConcentrationOfSpeciesA"); ca.setDepth(1);

### //! molar concentration of species B

@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 0, 1> cb; cb.setEntryName("MolarConcentrationOfSpeciesB"); cb.setDepth(1);

▶ @StateVariable (or equivalently @Output) declares new **scalar** state variables.



## //! molar concentration of species A

@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 0, 1> ca; ca.setEntryName("MolarConcentrationOfSpeciesA"); ca.setDepth(1);

### //! molar concentration of species B

@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 0, 1> cb; cb.setEntryName("MolarConcentrationOfSpeciesB"); cb.setDepth(1);

- @StateVariable (or equivalently @Output) declares new scalar state variables.
- ► The depth of a state variable x has the following meaning:
  - **–** 0 : only  $x|_{t+\Delta t}$ , is available and associated with a variable named x.
  - = 1 :  $x|_t$  is associated with the variable  $\mathbf{x}_{-}1$  and  $x|_{t+\Delta t}$  is associated with the variable  $\mathbf{x}_{-}$ .
  - higher values are hardly ever been used and highly discouraged.



### //! molar concentration of species A

@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 0, 1> ca; ca.setEntryName("MolarConcentrationOfSpeciesA"); ca.setDepth(1);

### //! molar concentration of species B

@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 0, 1> cb; cb.setEntryName("MolarConcentrationOfSpeciesB"); cb.setDepth(1);

- @StateVariable (or equivalently @Output) declares new scalar state variables.
- ightharpoonup The depth of a state variable x has the following meaning:
  - **–** 0 : only  $x|_{t+\Delta t'}$  is available and associated with a variable named x.
  - = 1 :  $x|_t$  is associated with the variable  $x_1$  and  $x|_{t+\Delta t}$  is associated with the variable x.
  - higher values are hardly ever been used and highly discouraged.
- ► The quantity type allows to define new quantities. Its integer template parameters have the following meaning in SI:

$$kg^{i_1} m^{i_2} s^{i_3} A^{i_4} K^{i_5} cd^{i_6} mol^{i_7}$$



### //! molar concentration of species A

@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 0, 1> ca; ca.setEntryName("MolarConcentrationOfSpeciesA"); ca.setDepth(1);

### //! molar concentration of species B

@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 0, 1> cb; cb.setEntryName("MolarConcentrationOfSpeciesB"); cb.setDepth(1);

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▶ Variables can be documented using a doxygen-like syntax.

# The @Function keyword

```
@Function ChemicalReaction {
    const auto B = k2 * (ca_1 + cb_1);
    const auto K = k1 + k2;
    const auto e = exp(- K * dt);
    ca = ca_1 * e + (B / K) * (1 - e);
    cb = ca_1 + cb_1 - ca;
}
```

- Several functions can be defined, although this feature has hardly been used in pratice. It is highly recommended to define only one function per model.
  - The '@Integrator' keyword has been introducted in Version 4.1 for consistency with behaviours and does not expect a function name;
- ▶ The implementation readily translates the analytical formula in C++ :

$$[A]|_{t+\Delta t} = \frac{B}{K} + \left(\frac{K [A]|_t - B}{K}\right) \exp(-K \Delta t)$$

with: 
$$B = k_2 ([A]|_t + [B]|_t)$$
 and  $K = k_1 + k_2$ .



# **Compilation and testing with MTest**

▶ mfront --obuild --interface=generic ChemicalReaction1.mfront



# **Compilation and testing with MTest**

- ▶ mfront --obuild --interface=generic ChemicalReaction1.mfront
- ▶ mtest ChemicalReaction1.mtest



# **Compilation and testing with MTest**

- ▶ mfront --obuild --interface=generic ChemicalReaction1.mfront
- mtest ChemicalReaction1.mtest
- ► MTest generates a file named ChemicalReaction1.res which contains the evolution of the state variables as a function of time.



# Results of the first test

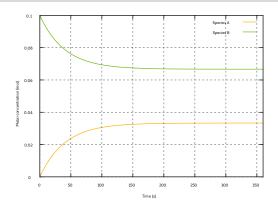


FIGURE – Evolution of the molar concentrations of species [A] and [B].

▶ tplot "ChemicalReaction1.res" -u 1:2 -title="Species
A" "ChemicalReaction1.res" -u 1:3 -title="Species
B" -with-grid -xlabel="Time (s)" -ylabel="Molar



# **Adding unit tests**

```
// loading the model
@Model 'src/libModel.so' 'ChemicalReaction1':
// initial value of the molar concentration of species B
@Real 'B0' 0.1:
@StateVariable 'MolarConcentrationOfSpeciesB' 'B0':
// time discretization
@Times {
 0, 360 in 100
// some useful variables
@Real 'k1' 'ChemicalReaction1::ReactionRateCoefficientAB':
@Real 'k2' 'ChemicalReaction1::ReactionRateCoefficientBA':
@Real 'K' 'k1 + k2':
@Real 'B' 'k2 * B0';
// unit tests
@Test<function>'MolarConcentrationOfSpeciesA''(B/K) * (1 - \exp(-K * t))' 1e-14:
@Test<function>'MolarConcentrationOfSpeciesB' 'B0 - (B/K) * (1 - exp(-K * t))' 1e-14;
```

▶ Unit tests can be defined using analytical solutions or reference files.



# Using the MTest Python module

```
import std import fel. tests import mest. WerboseLevel. VERBOSE_QUIET)

m = mtest.MTest()
m.setAuthor("Thomas Helfer")
m.setDate("09/08/2022")
m.setModel("generic", 'src/libModel.so', 'ChemicalReaction1')
m.setStateVariableInitialValue("MolarConcentrationOfSpeciesB", 0.1)
m.setStateVariableInitialValue("MolarConcentrationOfSpeciesB", 0.1)
m.setTimes([3.6 * i for i in range(0, 100]))
output_file = "ChemicalReaction1—python.res".format(k1)
m.setOutputFileName(output_file)
m.exectUef)
```

► The MTest Python module can be used to identify models using the LDC-OD component.



# Parametric studies with the MTest Python module

```
import std
import tfel.tests
import mtest
mtest.setVerboseMode(mtest.VerboseLevel.VERBOSE QUIET)
0.004166666666666667, 0.033333333333333333333
         0.06666666666666667]:
   m = mtest, MTest()
   m.setAuthor("Thomas Helfer")
   m.setDate("09/08/2022")
   m.setModel('generic', 'src/libModel.so', 'ChemicalReaction1')
   m.setStateVariableInitialValue('MolarConcentrationOfSpeciesB', 0.1)
   m.setParameter('ReactionRateCoefficientAB', float(k1))
   m.setTimes([3.6 * i for i in range(0, 100)])
   output_file = "ChemicalReaction1 - python - parametric - {}.res".format(k1)
   m.setOutputFileName(output file)
   m.execute()
```

► See the MFront' book to see how to perform parametric studies in bash



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# Reaction rate coefficients as functions of the temperature

Material properties	Value
k <sub>10</sub>	0.018377505387559667 s <sup>-1</sup>
k <sub>20</sub>	0.01013198112809354 s <sup>-1</sup>
$T_{a1}$	3000 K <sup>-1</sup>
$T_{\alpha 2}$	1500 K <sup>-1</sup>

▶ The reaction rate coefficients  $k_1$  and  $k_2$  are now assumed to depend on the temperature following the Arrhenius law, as follows:

$$\begin{cases} k_1 = k_{10} \exp\left(-\frac{T}{T_{a1}}\right) \\ k_2 = k_{20} \exp\left(-\frac{T}{T_{a2}}\right) \end{cases}$$

where  $k_{10}$ ,  $T_{a1}$ ,  $k_{20}$  and  $T_{a2}$  are material coefficients.



# **Numerical schemes**

- No closed-form solution exists.
- ► The ordinary differential equation can be integrated over the time step as follows :

$$\Delta \vec{Y} = \int_{t}^{t+\theta \Delta t} \vec{G}\left(\vec{Y}(\tau), \vec{Z}(\tau)\right) d\tau$$

- ► Various numerical schemes can be built by approximating the integral on the right hand side. In this tutorial, we consider:
  - The trapezoidal rule
  - The generalized mid-point rule

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#### **Code factorization**

▶ All the implementations based on the Model DSL will share the same internal state variables, the same external state variable (the temperature) and the same parameters.



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- ► All the implementations based on the Model DSL will share the same internal state variables, the same external state variable (the temperature) and the same parameters.
- Moreover, the parameters' declaration can be shared between all DSLs.
- ▶ It is thus convenient to factorize their declaration in two auxiliary files :
  - ChemicalReaction-parameters.mfront which contains the declaration of the parameters. This file can be imported by all DSLs.
  - ChemicalReaction-common.mfront contains the declaration of the internal state variables and the external state variable and imports the ChemicalReaction-parameters.mfront file. This file is only compatible with the Model DSL.



#### **Code factorization**

- ➤ All the implementations based on the Model DSL will share the same internal state variables, the same external state variable (the temperature) and the same parameters.
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- ▶ It is thus convenient to factorize their declaration in two auxiliary files :
  - ChemicalReaction-parameters.mfront which contains the declaration of the parameters. This file can be imported by all DSLs.
  - ChemicalReaction-common.mfront contains the declaration of the internal state variables and the external state variable and imports the ChemicalReaction-parameters.mfront file. This file is only compatible with the Model DSL.
- ► The ChemicalReaction-common.mfront file can be imported by the @Import keyword, as follows:

@Import "ChemicalReaction - common.mfront";



#### The ChemicalReaction-common.mfront file

```
@UseQt true;
@UnitSystem SI;

//! molar concentration of species A
@StateVariable quantity-real, 0, 0, 0, 0, 0, 1 > ca;
ca.setEntryName("MolarConcentrationOfSpeciesA");
ca.setDepth(1);

//! molar concentration of species B
@StateVariable quantity-real, 0, 0, 0, 0, 0, 0, 1 > cb;
cb.setEntryName("MolarConcentrationOfSpeciesB");
cb.setDepth(1);

@ExternalStateVariable temperature T;
T.setGlossaryName("Temperature");
T.setDepth(1);

@Import "ChemicalReaction—parameters.mfront";
```



### The ChemicalReaction-parameters.mfront file

```
//! reference rate coefficient of the reaction transforming species A to species B
```

@Parameter frequency k01 = 0.018377505387559667;

k01.setEntryName("ReferenceReactionRateCoefficientAB");

//! reference rate coefficient of the reaction transforming species B to species A

@Parameter frequency k02 = 0.01013198112809354;

k02.setEntryName("ReferenceReactionRateCoefficientBA");

//! activation temperature reaction transforming species B to species A

@Parameter temperature Ta1 = 3000;

Ta1.setEntryName("ActivationTemperatureAB");

//! activation temperature reaction transforming species B to species A

@Parameter temperature Ta2 = 1500;

Ta2.setEntryName("ActivationTemperatureBA");



#### Trapezoidal rule

► The integral on the right hand size can be approximated by the trapezoidal rule as follows :

$$\Delta \vec{Y} \approx \frac{\Delta t}{2} \left( \vec{G} \left( \vec{Y} (t), \vec{Z} (t) \right) + \vec{G} \left( \vec{Y} (t + \Delta t), \vec{Z} (t + \Delta t) \right) \right)$$



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▶ Applied to the chemical reaction example, this scheme leads to the approximation of the increment  $\Delta$  [A]:

$$\Delta [A] = \frac{B}{1 + K}$$

where:

$$-B = \Delta t \left( \langle k_2 \rangle \left[ B \right] \right|_t - \langle k_1 \rangle \left[ A \right] \right|_t \right) \text{ with } \begin{cases} \langle k_1 \rangle = \frac{k_1 \left( T \right|_{t+\Delta t} \right) + k_1 \left( T \right|_t \right)}{2} \\ \langle k_2 \rangle = \frac{k_2 \left( T \right|_{t+\Delta t} \right) + k_2 \left( T \right|_t \right)}{2} \end{cases}$$

$$- K = \frac{\Delta t}{2} \left( k_1 \left( T|_{t+\Delta t} \right) + k_2 \left( T|_{t+\Delta t} \right) \right).$$



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$$\Delta [A] = \frac{B}{1 + K}$$

where:

$$-B = \Delta t \left( \langle k_2 \rangle \left[ B \right] |_t - \langle k_1 \rangle \left[ A \right] |_t \right) \text{ with } \begin{cases} \langle k_1 \rangle = \frac{k_1 \left( T |_{t+\Delta t} \right) + k_1 \left( T |_t \right)}{2} \\ \langle k_2 \rangle = \frac{k_2 \left( T |_{t+\Delta t} \right) + k_2 \left( T |_t \right)}{2} \end{cases}$$

$$- K = \frac{\Delta t}{2} \left( k_1 \left( T|_{t+\Delta t} \right) + k_2 \left( T|_{t+\Delta t} \right) \right).$$

▶ This closed-form expression of  $\Delta$  [A] is due to the linear nature of the function  $\vec{G}$ .



# Implementation using the trapezoidal rule

```
@DSL Model:
@Model ChemicalReaction2:
@Author Thomas Helfer;
@Date 09/07/2022:
@Import "ChemicalReaction - common.mfront";
@Function ChemicalReaction {
 constexpr auto zero = quantity<real, 0, 0, 0, 0, 0, 0, 1>{};
 const auto k1 bts = k01 * exp(-T 1 / Ta1);
 const auto k1 ets = k01 * exp(-T/Ta1):
 const auto k2_bts = k02 * exp(-T_1 / Ta2);
 const auto k2 \text{ ets} = k02 * \exp(-T / Ta2);
 const auto mean k1 = (k1 bts + k1 ets) / 2;
 const auto mean_k2 = (k2_bts + k2_ets) / 2;
 const auto B = dt * (mean k2 * cb 1 - mean k1 * ca 1);
 const auto K = dt * (k1 ets + k2 ets) / 2;
 ca = ca 1 + B / (1 + K):
 cb = ca 1 + cb 1 - ca:
 if (cb < zero) {
   cb = zero:
   ca = ca 1 + cb 1:
 if (ca < zero) {
   ca = zero:
   cb = ca 1 + cb 1;
```



#### The generalized midpoint rule

► The general ordinary differential equation can be integrated over the time step using the generalized midpoint rule as follows :

$$\Delta \vec{Y} \approx \Delta t \vec{G} \left( \vec{Y} \Big|_{t+\theta \Delta t}, \vec{Z} \Big|_{t+\theta \Delta t} \right)$$

where  $\theta$  is a numerical parameter between 0 and 1.



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where  $\theta$  is a numerical parameter between 0 and 1.

▶ Applied to the chemical reaction example, this scheme leads to the approximation of the increment  $\Delta$  [A]:

$$\Delta [A] = \frac{B}{1 + \theta K}$$

with 
$$\begin{cases} B = \Delta t \left( k_2 \left( T|_{t+\theta \Delta t} \right) \left[ B \right] |_t - k_1 \left( T|_{t+\theta \Delta t} \right) \left[ A \right] |_t \right) \\ K = \Delta t \ k_1 \left( T|_{t+\theta \Delta t} \right) \end{cases}$$



# Implementation using the midpoint rule

```
@DSL Model:
@Model ChemicalReaction3:
@Author Thomas Helfer:
@Date 09/07/2022:
@Import "ChemicalReaction—common.mfront":
//! numerical parameter of the generalized mid—point rule
@Parameter real theta = 0.5:
theta.setEntrvName("Theta"):
@Function ChemicalReaction {
 constexpr auto zero = quantity<real, 0, 0, 0, 0, 0, 0, 1>{};
 const auto T mts = T 1 * (1 - \text{theta}) + \text{theta} * T;
 const auto k1 mts = k01 * exp(-T mts / Ta1);
 const auto k2 mts = k02 * exp(-T mts / Ta2):
 const auto B = dt * (k2 mts * cb 1 - k1 mts * ca 1);
 const auto K = dt * (k1 mts + k2 mts);
 ca = ca 1 + B / (1 + K * theta):
 cb = ca 1 + cb 1 - ca:
 // imposing positivity of the molar concentrations
 if (cb < zero){
   cb = zero:
   ca = ca 1 + cb 1;
 if (ca < zero){
   ca = zero:
   cb = ca 1 + cb 1;
```



#### A third scheme

- ▶ If the temperature increment is assumed small over a time step, an interesting scheme is to resuse the first one and evaluate the rate coefficients  $k_1$  and  $k_2$  at the middle of the time step  $t + \frac{\Delta t}{2}$ .
- ► The advantage of this implementation is that the exact solution is retrieved if the temperature is constant.



### Implementation of the third scheme

```
@DSL Model;
@Model ChemicalReaction4;
@Author Thomas Helfer;
@Date 09/07/2022;

@Import "ChemicalReaction—common.mfront";

@Function ChemicalReaction {
    const auto T_mts = (T_1 + T) / 2;
    const auto k1_mts = k01 * exp(-T_mts / Ta1);
    const auto k2_mts = k02 * exp(-T_mts / Ta2);
    const auto B = k2_mts * (ca_1 + cb_1);
    const auto K = k1_mts + k2_mts;
    const auto e = exp(-K * dt);
    ca = ca_1 * e + (B / K) * (1 - e);
    cb = ca_1 + cb_1 - ca;
```

```
@Author Thomas Helfer;
@Date 09/08/2022;
@Model 'src/libModel.so' 'ChemicalReaction2';

@Real 'B0' 0.1;

@StateVariable 'MolarConcentrationOfSpeciesB' 'B0';

@Real T0' 700;
@Real T1' 400;
@Real 'tau0' 30;
@Exel 'tau0' 30;
@ExternalStateVariable<function> Temperature' T0 + T1 * sin(t/tau0)';

@Times {
0, 200 in 100, 720 in 20
};
```

► The temperature is a periodic function of time with a period shorter than the relaxation time of the chemical reaction.

The DefaultModel DSL



# First implementation

```
@DSL DefaultModel:
@Model ChemicalReaction5:
@Author Thomas Helfer:
@Date 09/07/2022:
@UseQt true;
@UnitSystem SI:
//! molar concentration of species A
@AuxiliaryStateVariable quantity<real, 0, 0, 0, 0, 0, 1> ca;
ca.setEntryName("MolarConcentrationOfSpeciesA");
//! molar concentration of species B
@AuxiliaryStateVariable quantity<real, 0, 0, 0, 0, 0, 1> cb;
cb.setEntrvName("MolarConcentrationOfSpeciesB");
@Import "ChemicalReaction - parameters.mfront";
@Integrator{
 const auto T mts = T + dT / 2;
 const auto k1 mts = k01 * exp(-T mts / Ta1);
 const auto k2 mts = k02 * exp(-T mts / Ta2):
 const auto B = k2 \text{ mts} * (ca + cb);
 const auto K = k1 mts + k2 mts;
 const auto e = \exp(-K * dt):
 const auto sum = ca + cb:
 ca = ca * e + (B / K) * (1 - e);
 cb = sum - ca:
```

### **Improvements**

► A priori time step scaling factor

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- ► A priori time step scaling factor
- ► A posterior time step scaling factor



### Oxidation of pipes of Zircaloy alloys

 Many models of the litterature describing the oxidation of pipes made of Zircaloy alloys at constant temperature takes the following form:

$$I_0^2\left(\tau\right) = K\left(T\right)\,\tau$$

where  $I_0$  is the oxidation length and  $\tau$  the time from the beginning of the experiment.



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▶ The function K(T) is generally choosen of the form :

$$K(T) = K_0 \exp\left(-\frac{E_a}{RT}\right)$$

where  $K_0$  is characteristic of the pipe considered (material and size),  $E_a$  is an activation energy ( $J.mol^{-1}$ ) and R is the perfect gas constant.



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► A model appliable in case of non constant temperature can be obtained by differentiating the previous equation with respect to time :

$$2I_0 \frac{\mathrm{d}I_0}{\mathrm{d}\tau} = K(T)$$



#### **Numerical scheme**

► The previous Ordinary Differential Equation can be integrated over a time step  $[t:t+\Delta t]$  as follows :

$$|I_0|_{t+\Delta t}^2 - |I_0|_t^2 = \int_t^{t+\Delta t} K(T(\tau)) \Delta \tau \approx K(T + \theta \Delta T) \Delta t,$$

► This leads to the following update scheme :

$$I_0|_{t+\Delta t} = \sqrt{I_0|_t^2 + K(T + \theta \Delta T) \Delta t}$$





#### The Runge-Kutta algorithms

Runge-Kutta algorithms estimates the solution at the end of the time step by the weighted sum of approximations  $\Delta \vec{Y}$  of the increment  $\Delta \vec{Y}$ 

as follows: 
$$\vec{Y}\Big|_{t+\Delta t} = \vec{Y}\Big|_t + \sum_{i=1}^n b_i \Delta \vec{\tilde{Y}}^{(i)}$$



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$$\vec{Y}\Big|_{t+\Delta t} = \vec{Y}\Big|_t + \sum_{i=1}^n b_i \Delta \vec{\tilde{Y}}^{(i)}$$

▶ The approximations  $\Delta \, \tilde{\vec{Y}}^{(i)}$  are given by successive evaluations of the rate function  $\vec{G}$ :

$$\begin{cases} \Delta \vec{\tilde{Y}}^{(1)} = \Delta t \vec{G} \left( \vec{\tilde{Y}}^{(1)}, \vec{Z} \Big|_{t} \right) & \text{with } \vec{\tilde{Y}}^{(1)} = \vec{Y} \Big|_{t} \\ \Delta \vec{\tilde{Y}}^{(2)} = \Delta t \vec{G} \left( \vec{\tilde{Y}}^{(2)}, \vec{Z} \Big|_{t+c_{2} \Delta t} \right) & \text{with } \vec{\tilde{Y}}^{(2)} = \vec{Y} \Big|_{t} + a_{21} \Delta \vec{\tilde{Y}}^{(1)} \\ \vdots \\ \Delta \vec{\tilde{Y}}^{(n)} = \Delta t \vec{G} \left( \vec{\tilde{Y}}^{(n)}, \vec{Z} \Big|_{t+c_{n} \Delta t} \right) & \text{with } \vec{\tilde{Y}}^{(n)} = \vec{Y} \Big|_{t} + a_{n1} \Delta \vec{\tilde{Y}}^{(1)} + \ldots + a_{n,n-1} \Delta \vec{\tilde{Y}}_{n-1} \end{cases}$$

with  $\vec{Z}\Big|_{t+c_i \Delta t} = \vec{Z}\Big|_t + c_j \Delta \vec{Z}$  being the estimates of  $\vec{Z}$  at time  $t + c_j \Delta t$ .



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Runge-Kutta algorithms estimates the solution at the end of the time step by the weighted sum of approximations  $\Delta \vec{Y}$  of the increment  $\Delta \vec{Y}$ as follows:  $\vec{Y}\Big|_{t+\Delta t} = \vec{Y}\Big|_t + \sum_{i=1}^{m} b_i \Delta \vec{\tilde{Y}}^{(i)}$ 

lacktriangle The approximations  $\Delta \, ec{ ilde{Y}}^{(i)}$  are given by successive evaluations of the rate function  $\vec{G}$ :

$$\begin{cases} \Delta \vec{Y}^{(1)} = \Delta t \vec{G} \left( \vec{Y}^{(1)}, \vec{Z} \Big|_t \right) & \text{with} \quad \vec{Y}^{(1)} = \vec{Y} \Big|_t \\ \Delta \vec{Y}^{(2)} = \Delta t \vec{G} \left( \vec{Y}^{(2)}, \vec{Z} \Big|_{t+c_2 \Delta t} \right) & \text{with} \quad \vec{Y}^{(2)} = \vec{Y} \Big|_t + a_{21} \Delta \vec{Y}^{(1)} \\ \vdots \\ \Delta \vec{Y}^{(n)} = \Delta t \vec{G} \left( \vec{Y}^{(n)}, \vec{Z} \Big|_{t+c_n \Delta t} \right) & \text{with} \quad \vec{Y}^{(n)} = \vec{Y} \Big|_t + a_{n1} \Delta \vec{Y}^{(1)} + \ldots + a_{n,n-1} \Delta \vec{Y}_{n-1} \end{cases}$$
with  $\vec{Z} \Big|_{t+c_n \Delta t} = \vec{Z} \Big|_t + c_j \Delta \vec{Z}$  being the estimates of  $\vec{Z}$  at time  $t + c_j \Delta t$ .

with  $\vec{Z}\Big|_{t+c_i \wedge t} = \vec{Z}\Big|_t + c_j \Delta \vec{Z}$  being the estimates of  $\vec{Z}$  at time  $t + c_j \Delta t$ .

▶ A particular algorithm is thus caractherized by n,  $a_{ii}$ ,  $b_i$  and  $c_i$ .



# A classical fourth order Runge-Kutta algorithm

► The classical fourth order Runge-Kutta is based on the following scheme :

$$\vec{Y}\Big|_{t+\Delta t} = \vec{Y}\Big|_t + \frac{1}{6} \left( \Delta \vec{\tilde{Y}}^{(1)} + 2 \Delta \vec{\tilde{Y}}^{(2)} + 2 \Delta \vec{\tilde{Y}}^{(3)} + \Delta \vec{\tilde{Y}}^{(4)} \right)$$

with:

$$\begin{cases} \Delta \vec{\tilde{Y}}^{(1)} = \Delta t \vec{G} \left( \vec{Y}^{(1)}, \vec{Z} \Big|_t \right) & \text{with } \vec{Y}^{(1)} = \vec{Y} \Big|_t \\ \Delta \vec{\tilde{Y}}^{(2)} = \Delta t \vec{G} \left( \vec{Y}^{(2)}, \vec{Z} \Big|_t + \frac{\Delta \vec{Z}}{2} \right) & \text{with } \vec{Y}^{(2)} = \vec{Y} \Big|_t + \frac{\Delta \vec{Y}^{(1)}}{2} \\ \Delta \vec{\tilde{Y}}^{(3)} = \Delta t \vec{G} \left( \vec{Y}^{(3)}, \vec{Z} \Big|_t + \frac{\Delta \vec{Z}}{2} \right) & \text{with } \vec{Y}^{(3)} = \vec{Y} \Big|_t + \frac{\Delta \vec{Y}^{(2)}}{2} \\ \Delta \vec{\tilde{Y}}_4 = \Delta t \vec{G} \left( \vec{Y}^{(4)}, \vec{Z} \Big|_{t+\Delta t} \right) & \text{with } \vec{Y}^{(4)} = \vec{Y} \Big|_t + \Delta \vec{Y}^{(3)} \end{cases}$$



#### **Corrector/predictor algorithms**

- Some algorithms allow to compute a lower order approximation for free.
- ► The difference between the higher order estimate and the lower order approximation is an estimate of the integration error :
  - automatic sub-stepping.



#### **Available algorithms**

- Without substepping :
  - euler, rk2, rk4
- ► With automatic substepping :
  - rk42, rk54, rkCastem



#### Conventions of the RungeKuttaModel DSL

- ▶ The @Derivative code block is used to compute the rate  $\vec{Y}$  of the state variables for a given estimate  $\vec{Y}^{(i)}$ :
- ➤ In the @Derivative code block, the following conventions holds:
  - **–** dy denotes the value of the rate of the variable *y*.
  - y denotes the value of  $y^{(i)}$ .



# **Implementation**

```
@DSL RungeKuttaModel;
@Model ChemicalReaction6:
@UseOt true:
@UnitSystem SI;
@Algorithm rk54:
@Epsilon 1e-14;
@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 1> ca:
ca.setEntryName("MolarConcentrationOfSpeciesA");
@AuxiliaryStateVariable quantity<real, 0, 0, 0, 0, 0, 1> cb;
cb.setEntrvName("MolarConcentrationOfSpeciesB");
@Import "ChemicalReaction - parameters.mfront";
@LocalVariable quantity<real, 0, 0, 0, 0, 0, 0, 1> sum:
@InitLocalVariables {
 sum = ca + cb:
@Derivative{
 const auto k1 = k01 * exp(-T / Ta1):
 const auto k2 = k02 * exp(-T / Ta2);
 dca = k2 * sum - (k1 + k2) * ca;
@UpdateAuxiliaryStateVariables {
 cb = sum - ca:
```

The ImplicitModel DSL



@DSL ImplicitModel;

### **Implementation**

```
@Model ChemicalReaction7:
@Author Thomas Helfer:
@Date 09 / 07 / 2022:
@UseQt true;
@UnitSystem SI:
@Epsilon 1e-14;
@Theta 0.5:
//! molar concentration of species A
@StateVariable quantity<real, 0, 0, 0, 0, 0, 0, 1> ca:
ca.setEntrvName("MolarConcentrationOfSpeciesA");
//! molar concentration of species B
@AuxiliaryStateVariable quantity<real, 0, 0, 0, 0, 0, 1> cb;
cb.setEntryName("MolarConcentrationOfSpeciesB");
@Import "ChemicalReaction - parameters.mfront";
//! sum of the molar concentrations of species A and B
@LocalVariable quantity<real, 0, 0, 0, 0, 0, 0, 1> sum;
/*!
* reaction rate coefficient of the reaction transforming species A to species
* B at the middle of the time step
 */
@LocalVariable frequency k1 mts:
/*!
* reaction rate coefficient of the reaction transforming species B to species
* A at the middle of the time step
 */
@LocalVariable frequency k2 mts;
```

**Applications** 

#### **Conclusions**



- ▶ The examples of this tutorial can be downloaded here :
  - https://github.com/thelfer/MFrontBookExamples



# Thank you for your attention. Time for discussion!

https://tfel.sourceforge.net
https://www.researchgate.net/project/TFEL-MFront
https://twitter.com/TFEL\_MFront
https://github.com/thelfer/

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