

Numerical Methods for the reactive transport : Global DAE approach

Souhila Sabit
SAGE-INRIA, RENNES

co-authors

Jocelyne Erhel (IRISA, Rennes, France)
Édouard Canot (IRISA, Rennes, France)

Sophia Antipolis, July 2012

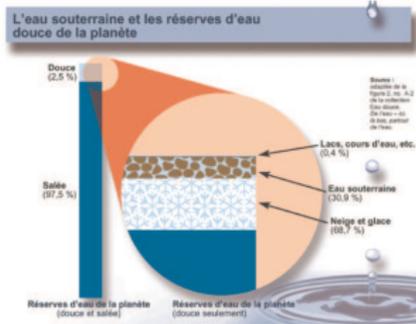
1 Introduction

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Introduction (1)

- **Modeling** reactive **transport** of contaminants in porous media is a complex problem combining the difficulties of **transport** modeling with those of modeling the **chemistry** and especially the **coupling** between the two.
- This modeling leads to a system of **partial differential equations and algebraic equations** whose unknowns are the quantities of chemical species.



S. Kräutle, P. Knabner, (2005); A new numerical reduction scheme for fully coupled multicomponent transport-reaction problems in porous media; Water Resources Research, Vol. 41, W09414, 17 pp.



S. Molins, J. Carrera, C. Ayora, Carlos and M.W. Saaltink, (2004); A formulation for decoupling components in reactive transport problems; Water Resources Research, Vol.40, W10301, 13 pp.

Introduction (2)

- Use method of lines, corresponding to the discretization in space only, leads to a differential algebraic system (DAE).
- Use a solver appropriate to solve this differential algebraic system (DAE).
- Use a solver based on an implicit method, in order and variable step, which requires at each time step, the iterative solution of large nonlinear system based on the Jacobian matrix
- Use Newton method to solve the nonlinear chemical systems.

our method **Global approach DAE**



C. de Dieuleveult, J. Erhel , M. Kern; A global strategy for solving reactive transport equations; Journal of Computational Physics,2009.

Chemical model (1)

- **Chemical reactions :**

$$\sum_{i=1}^{N_E} \nu_{r,i} E_i \rightleftharpoons 0, \quad r = 1, N_r \quad (1)$$

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- We assume that the system responses are independent so it is possible to reduce the system. We can express some unknown species which are called **secondary**, depending on other unknowns known as **components** .

- $E_\alpha \rightleftharpoons \sum_a \nu_{\alpha,a} E_a$
- $E_\sigma \rightleftharpoons \sum_a \nu_{\sigma,a} E_a + \sum_s \nu_{\sigma,s} E_s$
- $E_\pi \rightleftharpoons \sum_a \nu_{\pi,a} E_a$

Chemical model (2)

Chemical system

$$\begin{pmatrix} \mathbf{T}_a(x, t) - \mathcal{C}_a(\mathbf{l}c_a(x, t)) - \mathcal{F}_a(\mathbf{l}c_a(x, t), \mathbf{l}c_s(x, t), \mathbf{c}_\pi(x, t)) \\ \mathbf{T}_s(x, t) - \mathcal{F}_s(\mathbf{l}c_a(x, t), \mathbf{l}c_s(x, t)) \\ \mathcal{E}_\pi(\mathbf{l}c_a(x, t)) \end{pmatrix} = \mathbf{0} \quad (3)$$

Transport model (1)

The transport of these elements may be related to two phenomena, **advection** and **dispersion**.

- **Advection :**

$$-\nabla \cdot (c\mathbf{q}) = \mathcal{L}_{adv}(c) \quad (4)$$

- **Dispersion :**

$$\nabla \cdot (\mathbf{D}\nabla c) = \mathcal{L}_{disp}(c) \quad (5)$$

- **Transport equation :**

$$\varepsilon \frac{\partial c}{\partial t} = \mathcal{L}_{adv}(c) + \mathcal{L}_{disp}(c) = \mathcal{L}(c) \quad (6)$$

Coupling between chemistry and transport

- From the equations defined above, we obtain the following system:

$$\left\{ \begin{array}{l} \varepsilon \frac{\partial \mathbf{T}_a}{\partial t} = \mathcal{L}(\mathbf{C}_a) - \mathbf{f}, \\ \Phi(\mathbf{T}_a, \mathbf{y}) = \mathbf{0} \\ \mathbf{C}_a - \mathcal{C}_a(\mathbf{y}) = \mathbf{0} \\ + \text{boundary conditions} \\ + \text{initial conditions} \end{array} \right. \quad (7)$$

Where :

- The vector $\mathbf{y} = \begin{pmatrix} \mathbf{Ic}_a \\ \mathbf{Ic}_s \\ \mathbf{c}_\pi \end{pmatrix}$ corresponds to the unknown chemical system.
- Φ is the function describing the chemical phenomena defined in the system (3).

The global approach DAE

- We will write the **coupled** model as a **differential-algebraic** system and solve it by an appropriate solver after space discretization.

- The system (7) is written as:

$$\begin{cases} F(t, \mathbf{Y}, \frac{d\mathbf{Y}}{dt}) = \mathbf{0} \\ + \text{initials conditions at } t=0 \end{cases} \quad (8)$$

- with

$$\mathbf{Y} = \begin{pmatrix} \mathbf{T}_a \\ \mathbf{y} \\ \mathbf{C}_a \end{pmatrix} \text{ and } F \left(t, \mathbf{Y}, \frac{d\mathbf{Y}}{dt} \right) = \begin{pmatrix} \varepsilon \frac{d\mathbf{T}_a}{dt} - L(\mathbf{C}_a) + \mathbf{f} \\ \Phi(\mathbf{T}_a, \mathbf{y}) \\ \mathbf{C}_a - \mathcal{C}_a(\mathbf{y}) \end{pmatrix} \quad (9)$$

- Where \mathbf{T}_a , \mathbf{y} and \mathbf{C}_a are the variables discretized in space of size $N_a \times N_m$, $(N_a + N_s + N_\pi) \times N_m$ and $N_a \times N_m$ respectively.

GRT3D

- We solve the system with the **IDA** solver (**IDA** is a package that solves the Sundials algebraic differential equation (**DAE**) written as **F (t, y, y')= 0**).
- To solve the **chemical system** at the **initial** time **t=0**, we used **Kinsol** (**Kinsol** is a **nonlinear** solver for systems of algebraic equations based on **Newton's** algorithm).
- We provide two calculated functions to the solver, the value of **F** and its the **generalized Jacobian matrix** $J = \frac{\partial F}{\partial \mathbf{Y}} + \alpha \frac{\partial F}{\partial \mathbf{Y}'}$.
- This matrix is written as:

$$J = \begin{pmatrix} \alpha \mathbf{M} & \mathbf{0} & -\mathbf{L} \\ \frac{\partial \Phi}{\partial \mathbf{T}_a} & \frac{\partial \Phi}{\partial \mathbf{y}} & \mathbf{0} \\ \mathbf{0} & \frac{\partial C_a}{\partial \mathbf{y}} & \mathbf{I}_{N_m N_a} \end{pmatrix} \quad (10)$$

- **M** contains the discretized values of porosity
- α is a term from the time discretization
- **L** is the matrix of spatial discretization of the transport operator.
- The second block of rows contains the chemistry equations
- the third block of rows contains additional chemical equations.



GRT3D (2)

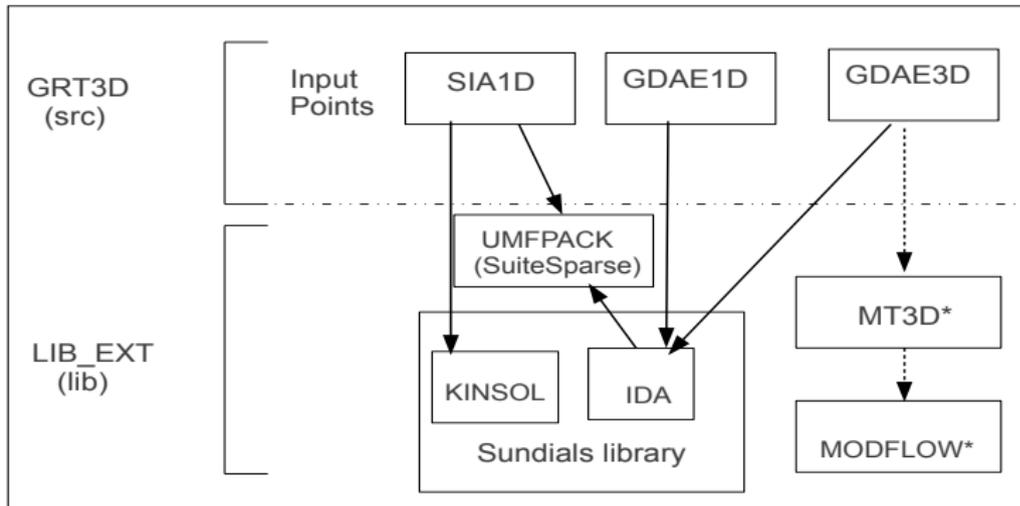


Figure: Description of GRT3D softwar

GRT3DR

- The Jacobian matrix (10) has a particular structure with **identity blocks** and **zero blocks**. We then **reduce** the size of the linear system at each Newton iteration.

- The system

$$J\mathbf{x} = \mathbf{g}$$

is equivalent to:

- $$\left\{ \begin{array}{l} \left(\frac{\partial \Phi}{\partial \mathbf{y}} - \frac{\partial \Phi}{\partial \mathbf{T}_a} \frac{\mathbf{M}^{-1}}{\alpha} \mathbf{L} \frac{\partial \mathcal{C}_a}{\partial \mathbf{y}} \right) \mathbf{x}_2 = \mathbf{g}_2 - \frac{\partial \Phi}{\partial \mathbf{T}_a} \frac{\mathbf{M}^{-1}}{\alpha} (\mathbf{g}_1 + \mathbf{L}\mathbf{g}_3), \\ \mathbf{x}_3 = \mathbf{g}_3 - \frac{\partial \mathcal{C}_a}{\partial \mathbf{y}} \mathbf{x}_2, \\ \mathbf{x}_1 = \frac{\mathbf{M}^{-1}}{\alpha} (\mathbf{g}_1 + \mathbf{L}\mathbf{x}_3), \end{array} \right. \quad (11)$$

- When $\mathbf{x} = (\mathbf{x}_1^T, \mathbf{x}_2^T, \mathbf{x}_3^T)^T$ and $\mathbf{g} = (\mathbf{g}_1^T, \mathbf{g}_2^T, \mathbf{g}_3^T)^T$. Thus solving the linear system is reduced to $(N_a + N_s + N_\pi) \times N_m$ instead of $(3N_a + N_s + N_\pi) \times N_m$.

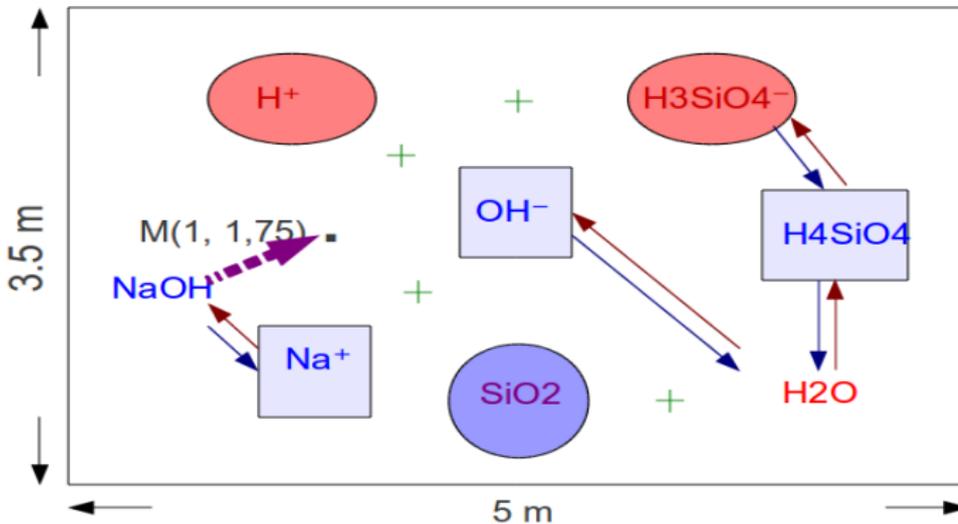
This method is implemented in **GRT3DR software**.

GRT3DR_SL

The use of the logarithm of the concentrations:

- We have the **product** of concentrations with constant **exponent** (2).
- We **linearize** all the equations
- The computation of **Jacobian matrix** is **very easy**.
- All the concentrations are **positive** (exponential).
- When concentrations are very small, the **Jacobian matrix** is **ill conditioned**.
- We have **zero** initial concentrations,, we approach the value by 10^{-10}
- We can also **remove** the **spectator ion** of the Chemical system.
- We use concentrations **without logarithm**, (Sans Log) then, we have the software **GRT3DR-SL** ,

test case Andra 2D (1)



- $\varepsilon = 1$ and $T_{final} = 1000$ days .



Mugler, G. and Bernard-Michel, G. and Faucher, G. and Miguez, R. and Gaombalet, J. and Loth, L. and Chavant, C.), Projet ALLIANCES: plan de qualification ; CEA, ANDRA, EDF.

test case Andra 2D (2)

- Initial values of T and logarithm

	Na^+	OH^-	H_4SiO_4
Initial values of T	0	0	10
log K	-32.2362	9.67086	8.28931

- The stoichiometric coefficient matrices

	Na^+	OH^-	H_4SiO_4	quartz
H^+	0	-1	0	0
$H_3SiO_4^-$	0	1	1	0
quartz	0	0	0	1

CPU time for the test Andra 2D (1)

- CPU time for the test Andra 2D with GRT3D

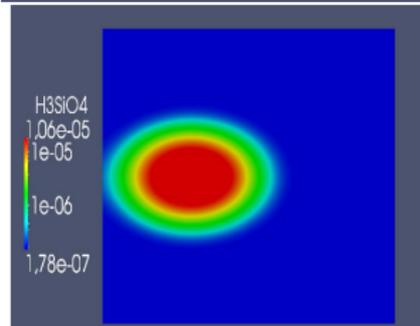
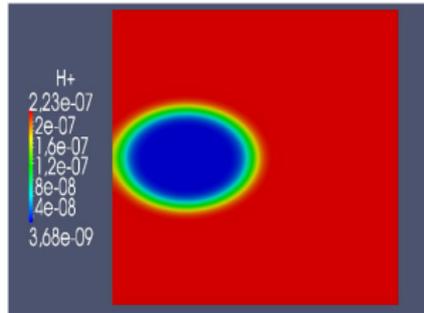
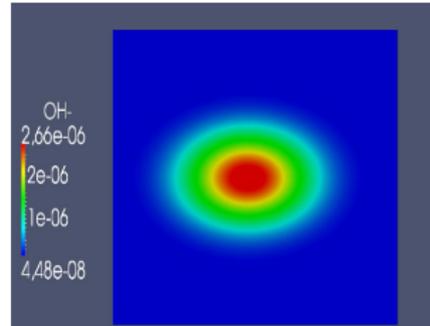
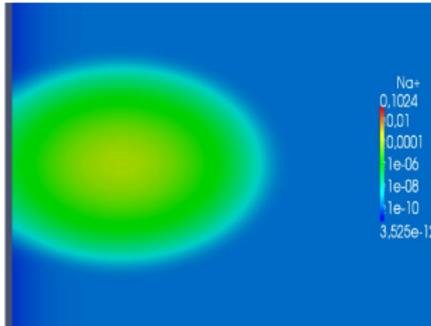
Mesh	GRT3D with spectator ion		GRT3D without spectator ion	
	size	CPU	size	CPU
21x14	2940	1 min 50 s	2058	18 s
48x28	13440	8 min	9408	1 min 36 s
81x56	45360	56 min	31752	6 min 33 s
71x101	71710	1h 20 min	50197	11 min 55 s
161x112	180320	3h	126224	32 min 43 s
322x224	721280	(1)	504896	(1)

- CPU time for the test Andra 2D without spectator ion for GRT3DR and GRT3DR-SL

Mesh	GRT3DR	GRT3DR-SL	Gain/GRT3D		size
21x14	4 s	2 s	0.5	0.11	882
48x28	21 s	8s	0.22	0.084	4032
81x56	1 min 53 s	50 s	0.29	0.13	13608
71x101	3 min 28 s	1 min 21 s	0.29	0.11	21513
161x112	16 min 30 s	4 min 32 s	0.5	0.14	54096
322x224	1 h 52 min	37 min 38 s			216384

(1): initial chemical balance not found

CPU time for the test Andra 2D (2)



Conclusion & perspectives

(a) Conclusion

- removing the spectator ion \Rightarrow reduce the CPU time by a factor of six
- GRT3DR \Rightarrow gain a factor two CPU time with respect to the CPU time of GRT3D
- GRT3DR-SL \Rightarrow reduce the CPU time by a factor between two and four compared to the CPU time of GRT3DR.

(b) Perspectives

- apply the SNIA method (Non-iterative sequential methods) and compare the CPU time.
- study the nonlinear complementarity problem.
- use semi-smooth Newton method.
- Parallelism through the components or the mesh.

Thank

Thank you for your attention