



A comparison between the second-order MOOD and MUSCL methods for the 1D shallow water problem in the presence of dry/wet interfaces

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- Presentation outline

- Motivation
- Generic second-order FV scheme
- MOOD vs. MUSCL
- Numerical results
- Conclusions and future work

- Motivation

- The SW problem with varying bathymetry has a **wide number of applications**: tsunami, flooding, coastal erosion...
- A large number of numerical schemes exists, in particular very high-order FV schemes.
- Still, **new 2nd-order methods are interesting** because these **techniques are widely used in engineering and environmental applications** due to their simplicity and computational efficiency **but need to be improved**.
- The **MUSCL** method is often used in this context, but recently a new method, **MOOD**, involving a **different limiting approach** has been proposed and tested, for example, for the Euler system and for the 2D-SW problem with varying bathymetry (up to 6th-order of accuracy on unstructured meshes).
- MOOD and MUSCL differ essentially in the **limiting procedure** used to prevent oscillations in the vicinity of discontinuities (*a priori* for **MUSCL**, *a posteriori* for **MOOD**).

- Motivation

- **Aim:** compare the “performance” of MOOD vs. MUSCL for the 1D-SW system with varying bathymetry.

Assess: { accuracy (order of convergence for smooth solutions)
shock capturing ability (sharpness and oscillations)
behaviour of the numerical solution near dry/wet interfaces



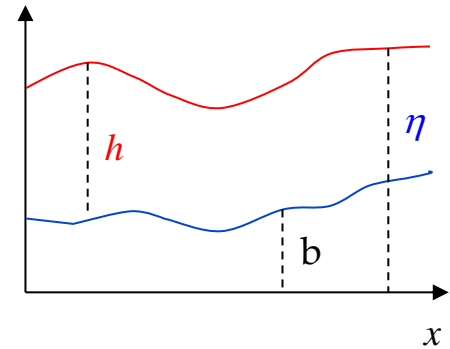
set of numerical simulations of different nature

- Generic 2nd-order FV scheme

❖ Non-linear 1D-SW system

$$\begin{cases} \partial_t h + \partial_x (hu) = 0, \\ \partial_t (hu) + \partial_x \left(hu^2 + \frac{g}{2} h^2 \right) = -gh \partial_x b, \end{cases}$$

hydrostatic condition: $\eta = h + b$



❖ Discretisation: time

$$[0, T] \rightarrow 0 = t^0 < t^1 < \dots < t^n < \dots < t^N = T$$

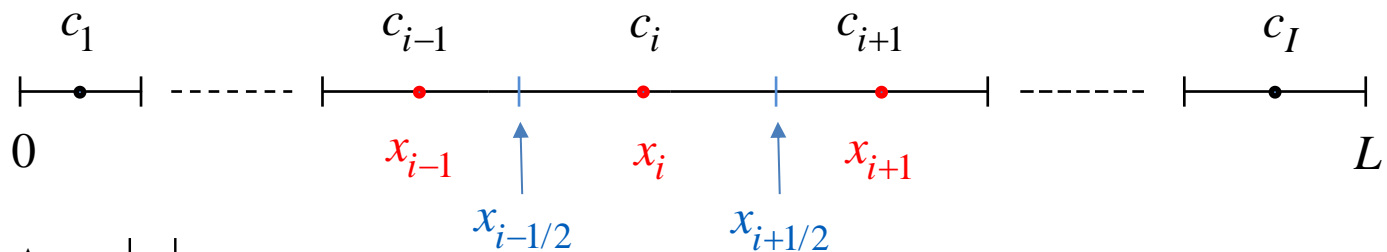
time step:

$$\Delta t_k = t^k - t^{k-1}, \quad k = 1, \dots, N. \quad \Delta t_k = \Delta t = \frac{T}{N} \quad \text{if regular subdivision}$$

- Generic 2nd-order FV scheme

❖ Discretisation: space

$\Omega = [0, L] \rightarrow I$ non-overlapping cells:



$$\Delta x_i = |c_i|$$

For a function ϕ at t^n : $\phi_i^n \approx \phi(x_i)$

$$\left\{ \begin{array}{l} \phi_{i+1/2,L}^n \approx \phi(x_{i+1/2}^-) \\ \phi_{i+1/2,R}^n \approx \phi(x_{i+1/2}^+) \end{array} \right.$$

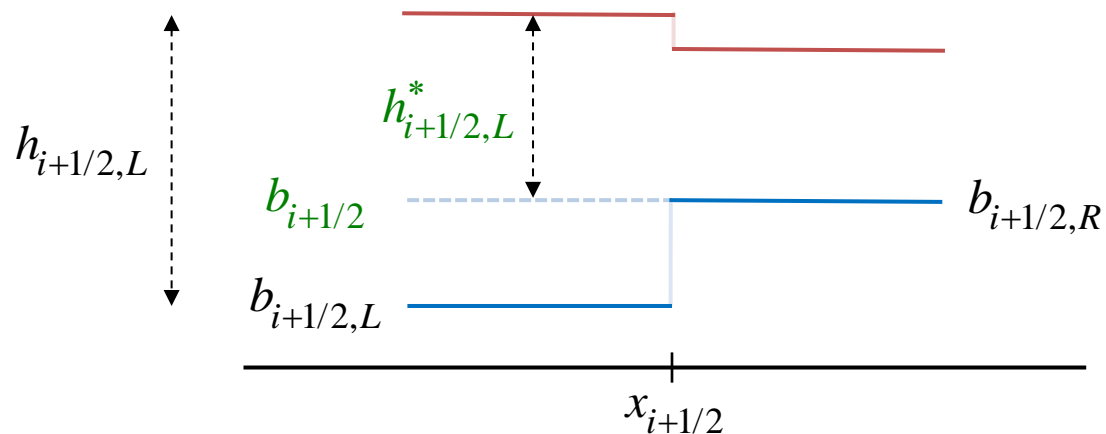
- Generic 2nd-order FV scheme

❖ **Hydrostatic reconstruction** (Audusse et al., 2004)

Define: $b_{i+1/2} = \max(b_{i+1/2,L}, b_{i+1/2,R})$

Set: $h_{i+1/2,K}^* = \max(0, h_{i+1/2,K} + b_{i+1/2,K} - b_{i+1/2})$,

$\eta_{i+1/2,K}^* = h_{i+1/2,K}^* + b_{i+1/2}$, $u_{i+1/2,K}^* = u_{i+1/2,K}$, with $K = L, R$.



- Generic 2nd-order FV scheme

- ❖ Generic second-order scheme (Audusse et al. methodology)

Setting $U = (h, hu, b)$, the discretisation of the 1D-SW system leads to:

$$U_i^{n+1} = U_i^n - \frac{\Delta t}{\Delta x_i} \left(\mathcal{F}_{i+1/2}^n + \varepsilon_{i+1/2,L}^n - \mathcal{F}_{i-1/2}^n - \varepsilon_{i-1/2,R}^n \right) - \Delta t S_i^n,$$

where

$$\mathcal{F}_{i-1/2}^n = F \left(U_{i-1/2,L}^{*,n}, U_{i-1/2,R}^{*,n} \right),$$

$$\varepsilon_{i-1/2,K}^n = \frac{g}{2} \left[(h_{i-1/2,K}^{*,n})^2 - (h_{i-1/2,K}^n)^2 \right], \quad \text{with } K = L, R,$$

$$S_i^n = -g \frac{h_{i+1/2,L}^n + h_{i-1/2,R}^n}{2} \frac{b_{i+1/2,L}^n - b_{i-1/2,R}^n}{\Delta x_i}.$$

- MOOD vs. MUSCL

➤ Local second-order reconstruction

Local linear reconstruction required → approximation of first derivatives:

$$\text{interfaces: } p_{i-1/2}(\phi) = \frac{\phi_i - \phi_{i-1}}{\Delta x}, \quad p_{i+1/2}(\phi) = \frac{\phi_{i+1} - \phi_i}{\Delta x}.$$

$$\text{centroid: } p_i(\phi) = \frac{\phi_{i+1} - \phi_{i-1}}{2\Delta x}.$$

But non-limited linear reconstruction will lead to **oscillations in the vicinity of discontinuities** and even in less extreme cases.



Need to **implement limiting procedures** → loss of accuracy!

- MOOD vs. MUSCL

➤ Limitation strategy

MUSCL (Monotonic Upstream-Centred Scheme for Conservation Laws; van Leer 1979):

slopes are reduced in order to verify some stability criterion;
corrections to slopes are made before computing the solution
(*a priori*).

MOOD (Multi-dimensional Optimal Order Detection; Clain, Diot, Loubère 2011):

assumes solution is smooth enough and the candidate solution is
computed without changing the slopes; corrections to slopes are
made if the candidate solution is “problematic” (*a posteriori*).



MOOD generally less intrusive, providing better accuracy

- MOOD vs. MUSCL

➤ MUSCL in brief

Reconstructed values at the left- and right-hand side of interfaces:

$$\phi_{i-1/2,R} = \phi_i - q_i(\phi) \frac{\Delta x}{2}, \quad \phi_{i+1/2,L} = \phi_i + q_i(\phi) \frac{\Delta x}{2}, \quad \text{with } \phi = \eta, h, hu,$$

$$b_{i-1/2,R} = \eta_{i-1/2,R} - h_{i-1/2,R}, \quad b_{i+1/2,L} = \eta_{i+1/2,L} - h_{i+1/2,L},$$

where

$$q_i(\phi) = \theta \left(p_{i-1/2,R}(\phi), p_{i+1/2,L}(\phi) \right)$$

van-Alabada

min-mod

van-Leer

- MOOD vs. MUSCL

➤ MOOD in brief

- Key concepts: Cell/Edge Polynomial Degree (CPD/EPD)

EPD → polynomial degree used to evaluate the reconstruction at both sides of each edge:

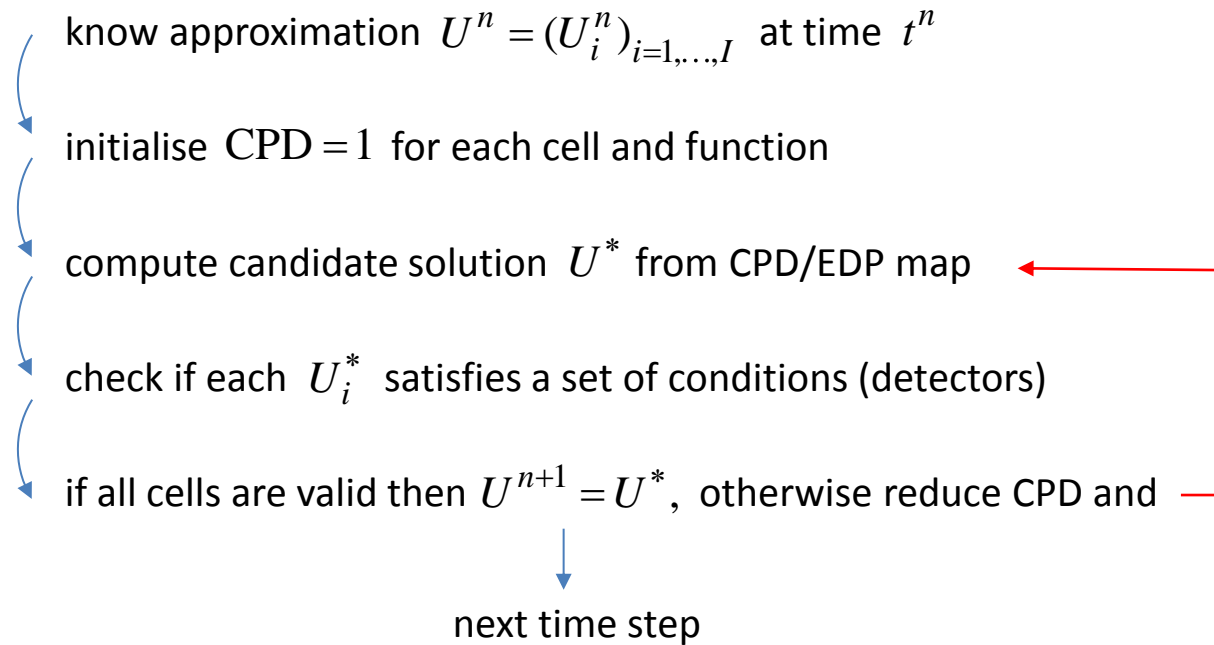
$$\text{EPD}_{i+1/2}(\phi) = \min(\text{CPD}_i(\phi), \text{CPD}_{i+1}(\phi))$$

$$\text{CPD}_i(\phi) = \begin{cases} 1 & \rightarrow \text{full slope used (second-order)} \\ 0 & \rightarrow \text{null slope used (first-order)} \end{cases}$$

- MOOD vs. MUSCL

➤ MOOD in brief

- MOOD loop:



- MOOD vs. MUSCL

➤ MOOD basic detectors

- **Aim:** determine if U_i^* is **eligible** (physically ok, no spurious oscillations)

Physical Admissible Detector: $h_i^* \geq 0$

(PAD)

Maximum Principle Detector: $\min(\phi_{i-1}^n, \phi_i^n, \phi_{i+1}^n) \leq \phi_i^* \leq \max(\phi_{i-1}^n, \phi_i^n, \phi_{i+1}^n)$

(MPD)

Extrema Detector: $\min(\phi_{i-1}^*, \phi_{i+1}^*) \leq \phi_i^* \leq \max(\phi_{i-1}^*, \phi_{i+1}^*)$

(ED)

- MOOD vs. MUSCL

➤ MOOD relaxation detectors

- **Aim:** distinguish real (smooth) extrema from extrema caused by local spurious oscillations (Gibbs phenomenon)

Local curvature indicator

$$C_i(\gamma) = \frac{\gamma_{i-1} - 2\gamma_i + \gamma_{i+1}}{(\Delta x)^2}, \quad i = 2, \dots, I-1; \quad C_1(\gamma) = C_2(\gamma), \quad C_I(\gamma) = C_{I-1}(\gamma).$$

$$\chi_{\min,i} = \min(C_{i-1}, C_i, C_{i+1}), \quad \chi_{\max,i} = \max(C_{i-1}, C_i, C_{i+1}), \quad i = 2, \dots, I-1.$$

$$\chi_{\min,i} = \chi_{\min,i}(\phi^*), \quad \chi_{\max,i} = \chi_{\max,i}(\phi^*).$$

- MOOD vs. MUSCL

➤ MOOD relaxation detectors

Small Curvature Detector: $\max(|\chi_{\min,i}|, |\chi_{\max,i}|) \leq \varepsilon_C.$

(SCD)

numerical solution is locally linear

Local Oscillation Detector: $\chi_{\min,i} \chi_{\max,i} \leq 0.$

(LOD)

local oscillation due to variation of curvature sign

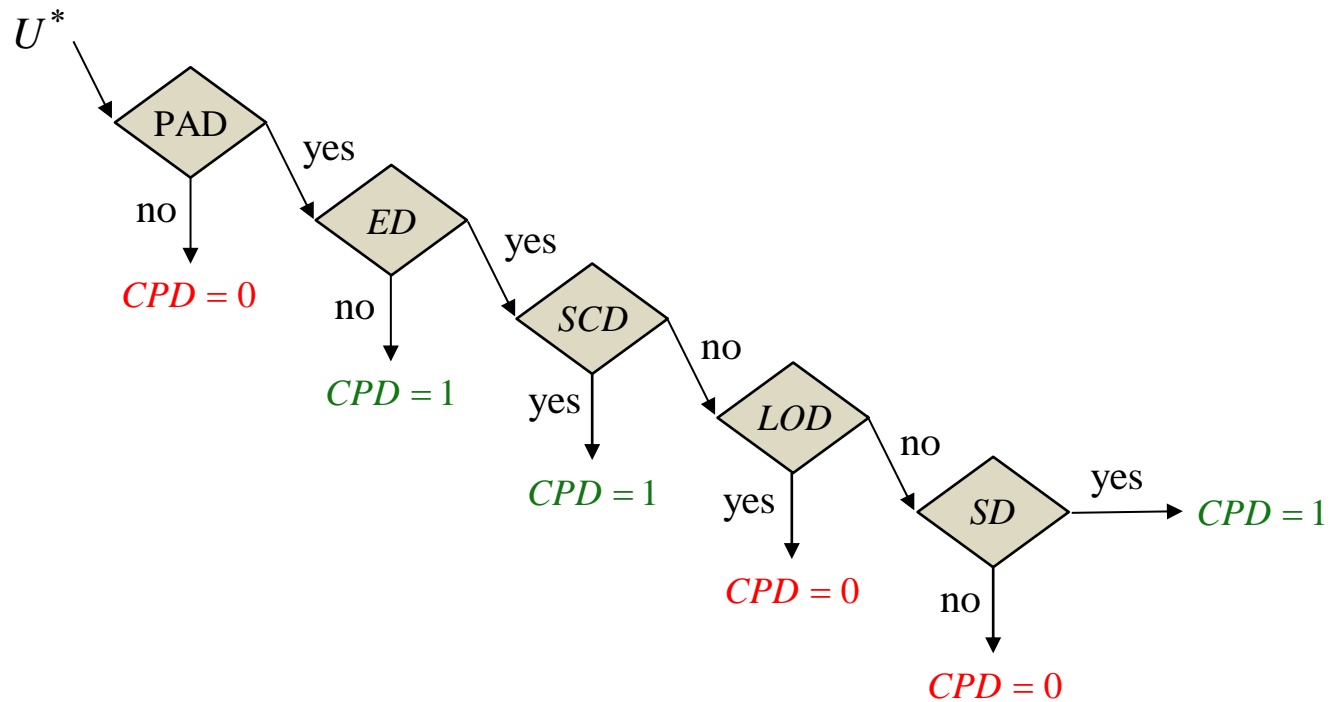
Smoothness Detector: $\frac{\min(|\chi_{\min,i}|, |\chi_{\max,i}|)}{\max(|\chi_{\min,i}|, |\chi_{\max,i}|)} \geq 1 - \varepsilon_S.$

(SD)

numerical solution is locally smooth

- MOOD vs. MUSCL

➤ MOOD detector algorithm



- Numerical results

□ General aspects

Time scheme: TVD-RK2, $CFL = 0.4$

Conservative flux scheme: HLL

Meshes: $|c_i| = \Delta x$

Dry/wet threshold: 10^{-6}

MUSCL limiter procedure: η, h, hu

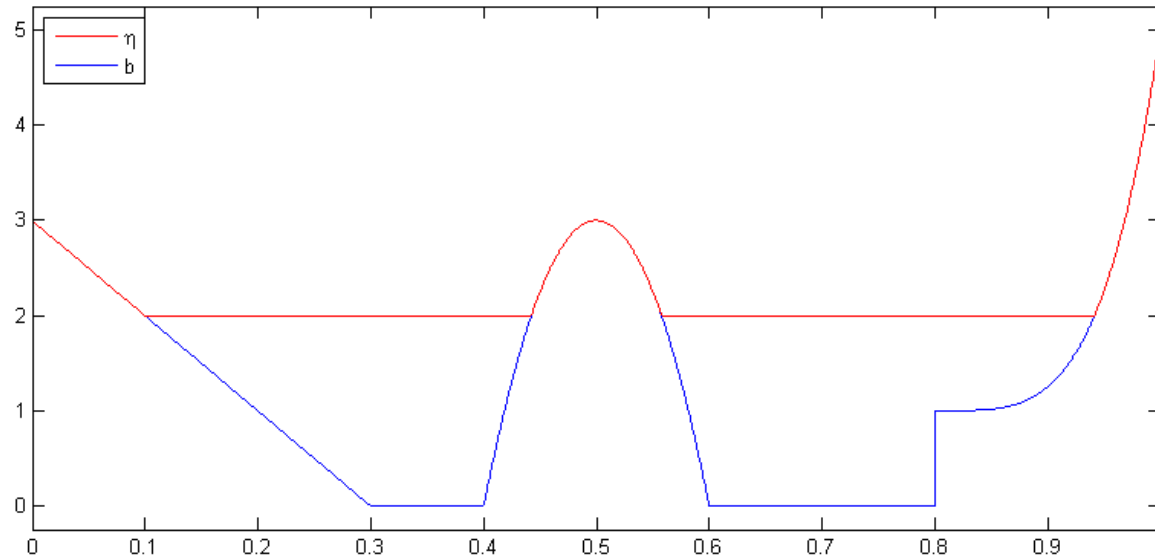
MOOD limiter procedure: h

$$\varepsilon_c = \delta^3, \text{ with } \delta = \Delta x/L; \quad \varepsilon_s = 0.5$$

Convergence assessment: L^1 - error: $\frac{1}{I} \sum_{i=1}^I |\phi_i^N - \phi_i^{ex}|$, L^∞ - error: $\max_i |\phi_i^N - \phi_i^{ex}|$

- Numerical results

☐ LAKE AT REST

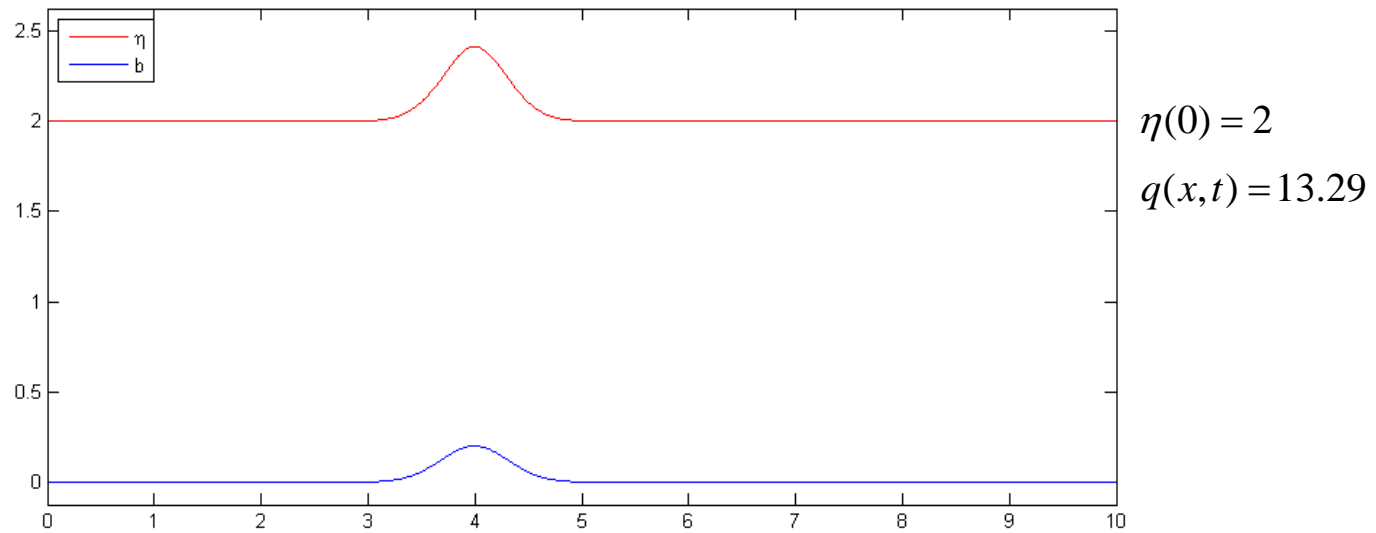


Meshes: 50, 100, 200 cells; $T = 1 \rightarrow$ up to 2215 time steps.

L^1 -error $< 10^{-15}$, L^∞ -error $< 10^{-14} \rightarrow$ C-property satisfied

- Numerical results

□ SMOOTH SOLUTION (steady-state supercritical flow)



Meshes: 100, ..., 1600 cells; $T = 10$; $b(x) = 0.2 \exp(-5(x-4)^2)$.

MUSCL detector only on h ; MOOD detector.

- Numerical results

SMOOTH SOLUTION (steady-state supercritical flow)

Nb of Cells	η				u			
	err_1		err_∞		err_1		err_∞	
100	2.44e-03	—	4.76e-02	—	5.40e-03	—	7.91e-02	—
200	5.20e-04	2.2	2.05e-02	1.2	1.12e-03	2.3	3.38e-02	1.2
400	1.12e-04	2.2	5.77e-03	1.8	2.38e-04	2.2	9.52e-03	1.8
800	2.30e-05	2.3	1.47e-03	2.0	4.89e-05	2.3	2.43e-03	2.0
1600	5.43e-06	2.1	3.73e-04	2.0	1.16e-05	2.1	6.15e-04	2.0

MUSCL

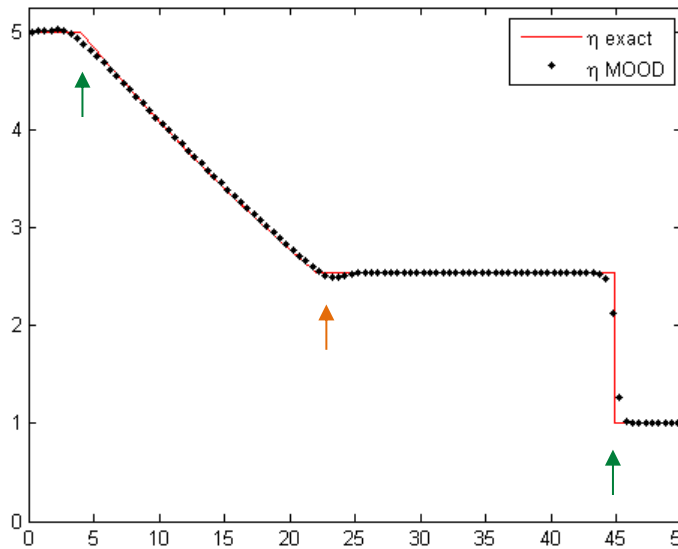
Nb of Cells	η				u			
	err_1		err_∞		err_1		err_∞	
100	1.91e-03	—	3.15e-02	—	5.06e-03	—	6.88e-02	—
200	5.23e-05	5.2	8.19e-04	5.3	1.24e-04	5.4	1.61e-03	5.4
400	8.49e-06	2.6	1.29e-04	2.7	2.13e-05	2.5	2.73e-04	2.6
800	1.45e-06	2.5	2.21e-05	2.5	3.81e-06	2.5	5.12e-05	2.4
1600	2.86e-07	2.3	4.26e-06	2.4	7.84e-07	2.3	1.07e-05	2.3

MOOD

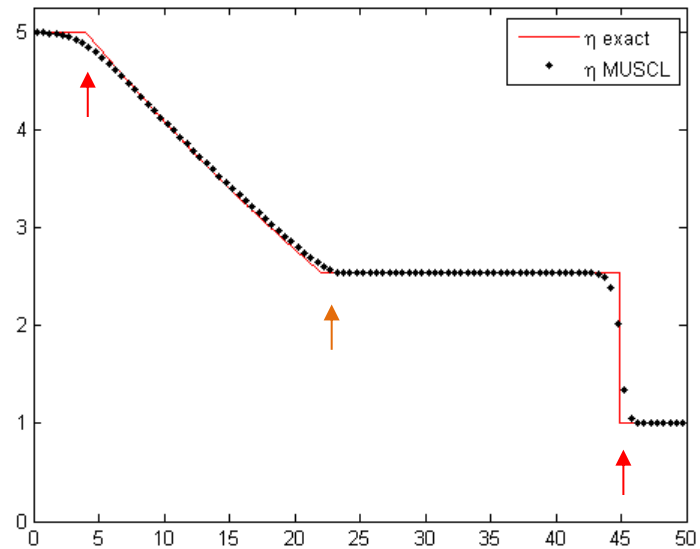
- Numerical results

□ DAM BREAK ON A WET BED

$$\text{Mesh: 100 cells; } T = 3; \quad \eta(x, 0) = \begin{cases} 5, & 0 \leq x \leq 25, \\ 1, & 25 \leq x \leq 50. \end{cases}$$



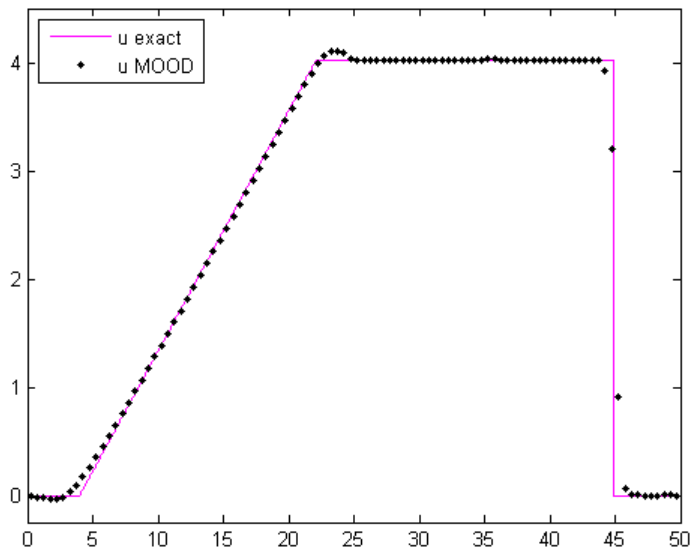
MOOD



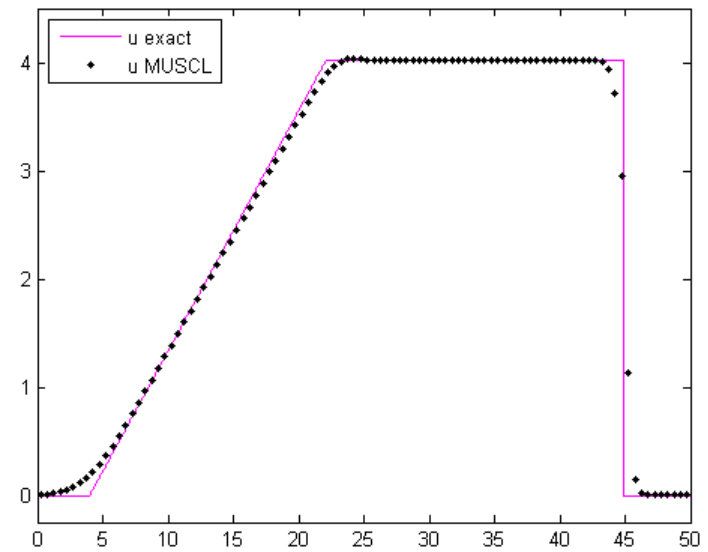
MUSCL

- Numerical results

DAM BREAK ON A WET BED



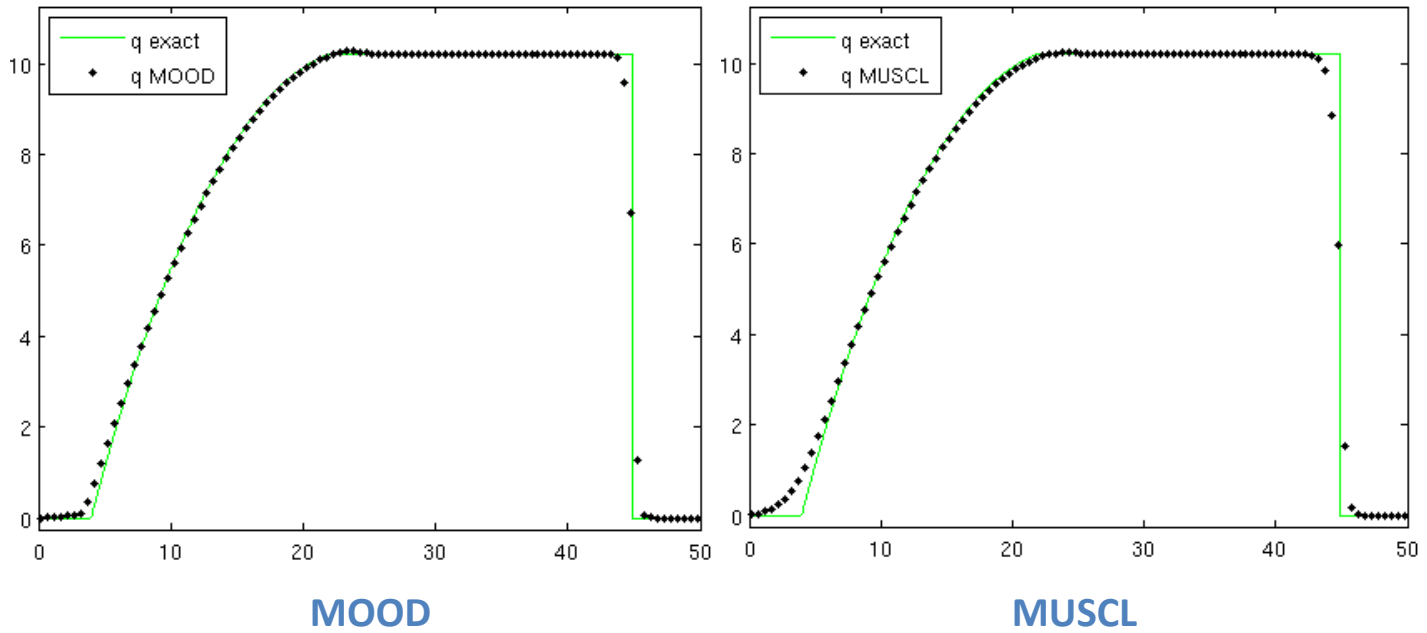
MOOD



MUSCL

- Numerical results

DAM BREAK ON A WET BED



- Numerical results

DAM BREAK ON A WET BED

Overall convergence:

$$\text{err}_1 = C (\Delta x)^\beta$$

MOOD:

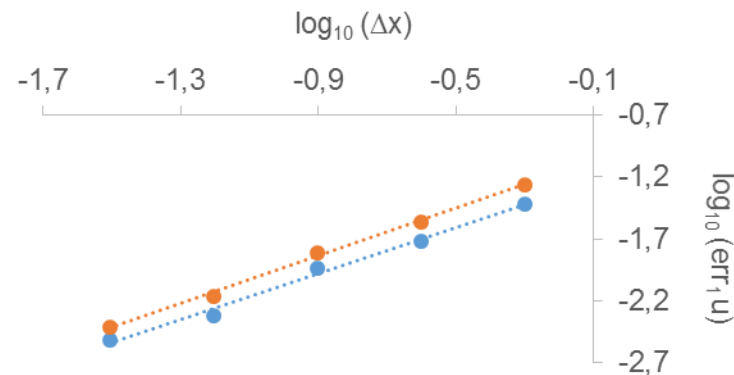
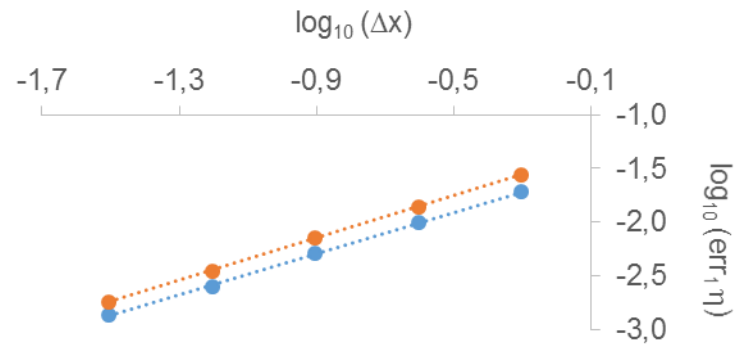
$$C_\eta = 0.037$$

$$C_u = 0.072$$

MUSCL:

$$C_\eta = 0.055$$

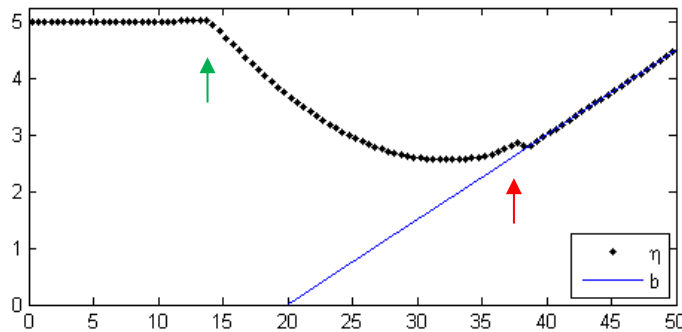
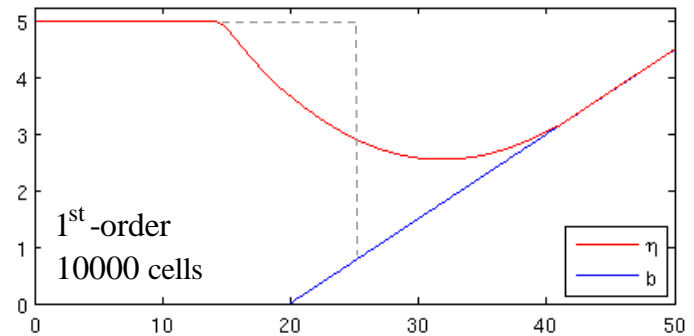
$$C_u = 0.107$$



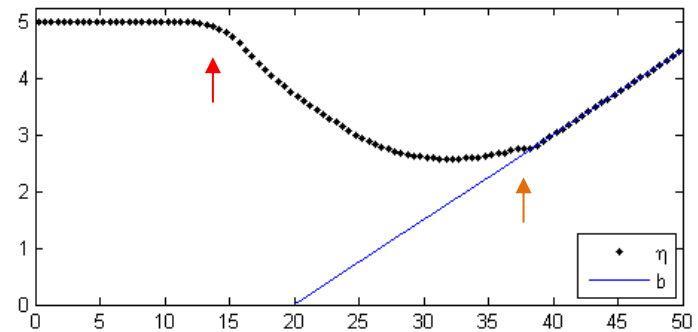
- Numerical results

□ DRY/WET SIMULATION (smooth bathymetry)

Mesh: 100 cells; $T = 1.5$.



MOOD

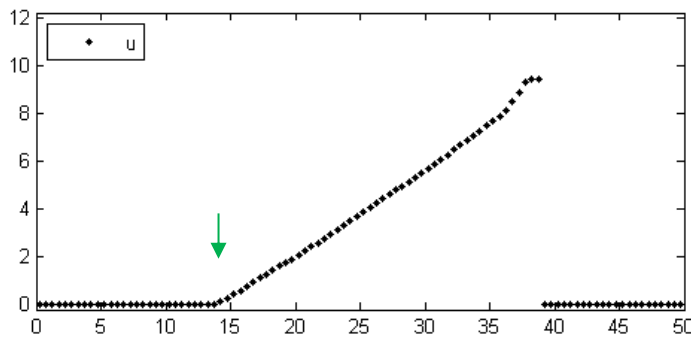
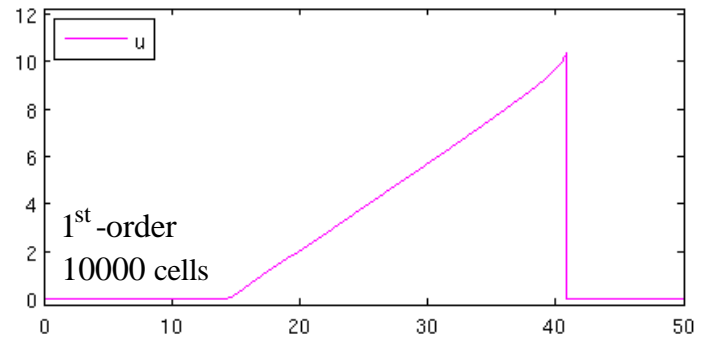


MUSCL

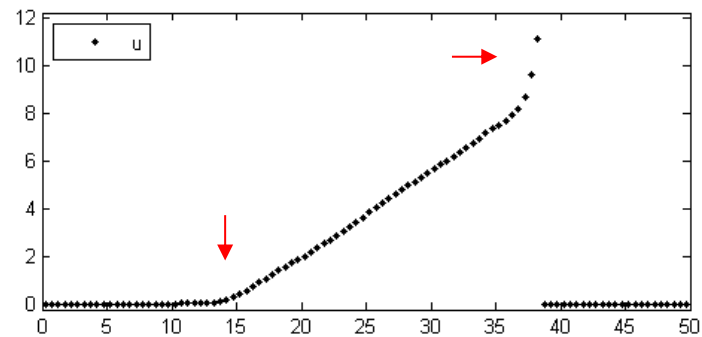
- Numerical results

DRY/WET SIMULATION (smooth bathymetry)

Mesh: 100 cells; $T = 1.5$.



MOOD



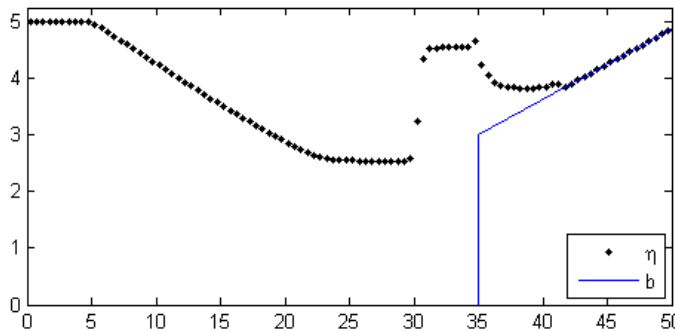
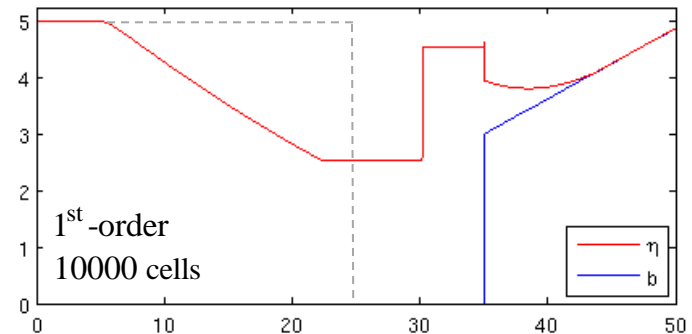
MUSCL

- Numerical results

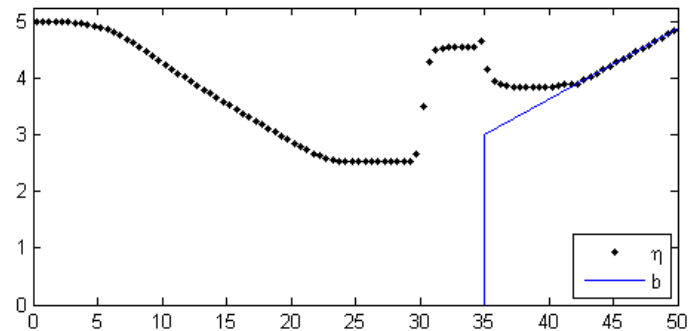
□ DRY/WET SIMULATION (discontinuous bathymetry)

Mesh: 100 cells; $T = 2.75$.

MOOD involves h and u !



MOOD

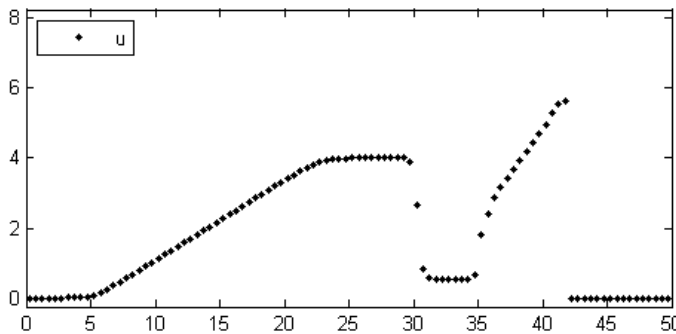
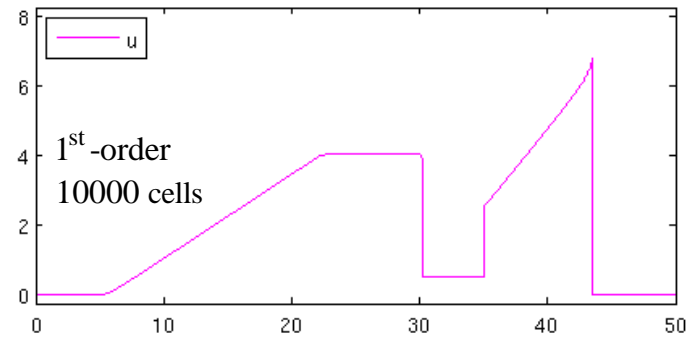


MUSCL

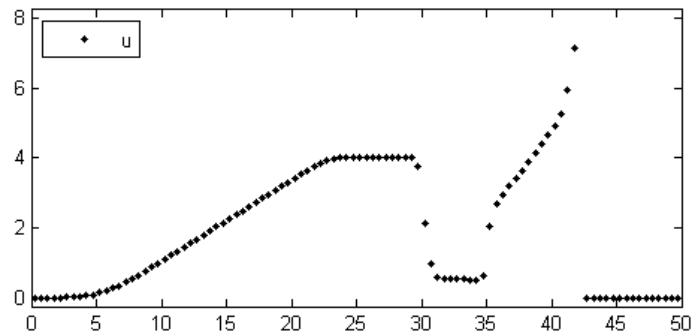
- Numerical results

DRY/WET SIMULATION (discontinuous bathymetry)

Mesh: 100 cells; $T = 2.75$.



MOOD



MUSCL

- Conclusions and future work

- We performed **comparisons** between the 2nd-order **MOOD** and the **MUSCL** methods for the **non-linear shallow water system with varying bathymetry**.
- Both methods comply with the C-property, but the **MOOD** method is **less diffusive** than the MUSCL method (shock, rarefaction propagation), and **is more accurate** when dealing with **smooth solutions**.
- **Dry/wet interfaces** with smooth/discontinuous bathymetry are reasonably **well handled by MOOD** technique (better velocity / kinetic energy estimates).
- Still, **new detectors** and **schemes** have to be proposed to improve the treatment of wet/dry interfaces (location, height and velocity profiles).