



### GLOFRIM

# Coupling of hydrologic and hydrodynamic models across scales for improved flood simulations

#### **Dirk Eilander**

E: dirk.eilander@vu.nl

T: @DirkEilander

W: www.compoundevents.org

#### Acknowledgements:

Jannis Hoch (University Utrecht) Hiroaki Ikeuchi (Unvirsity of Tokyo) Philip Ward (IVM, Amsterdam) Hessel Winsemius (TU Delft, Deltares)

### BACKGROUND

henkdeboer.nl

# **Compound Flooding**

#### Thailand floods: Bangkok 'impossible to protect'

3 20 October 2011 Asia-Pacific

😏 😒 🗹 < Share



#### **Bangladesh cyclone kills hundreds**

At least 600 people are reported to have died after a powerful cyclone smashed into Bangladesh's coast, levelling villages and uprooting trees.



When multiple flood drivers combine to affect flood impact

Jundrada of thousands



# (Global) Flood Modelling Status Quo



> Potentially underestimating risk

# (Global) Flood Modelling Compound Flooding



> Impact of combined flood drivers, taking into account dependence

### How to address compound floods?

Integration!

### > communities <u>www.compoundevent.org</u>

### > models Facilitating seamless coupling

### **CONCEPT AND APPLICATIONS**





### Integrated Modelling Framework //// /// //// meteorology temperature hydrology 🖌 precipitation precipitation runoff runoff routing (overbank) discharge

### Benchmark Routing – Amazon



Hoch, Eilander & Ikeuchi, in preparation <u>AGU poster</u>



### Global CaMa-Flood – GTSM Coupling



Eilander, in preparation AGU presentation

### Global CaMa-Flood – GTSM Coupling preliminary results

River mouth water level peaks



Eilander, in preparation AGU presentation

#### Integrated Modelling Framework //// /// //// meteorology temperature hydrology 🛓 precipitation precipitation runoff runoff routing (overbank) discharge full 2D floodplain hydrodynamics nearshore water levels offshore + nearshore hydrodynamics surge & tides waves

### Integrated Modelling Framework







### **GLOFRIM**

# Basic Model Interface (BMI)



API for component based integration of models

```
1
      # initialize model
      bmi.initialize_model()
 2
 3
      # get variable (numpy array)
 4
 5
      bmi.get_value('variable_name')
 6
 7
      # set variables
      bmi.set_value('variable_name', variable)
 8
      bmi.set_value_at_indices('variable_name', [1, 2, 23], variable)
 9
10
11
      # run models for one time step
      bmi.update()
12
                                             A component-based approach to integrated modeling in the
13
                                             geosciences: The design of CSDMS
      # finalize model
14
                                             Scott D. Peckham<sup>a</sup>, Eric W.H. Hutton<sup>a,*</sup>, Boyana Norris<sup>b</sup>
      bmi.finalize()
15
                                             https://github.com/csdms/bmi
16
```

# GLOFRIM v1.0

> Python scripts based on BMI

>> for specific hydrological and hydrodynamic models

>> Spatially explicit

>> Online coupling

GLOFRIM v1.0 – A globally applicable computational framework for integrated hydrological-hydrodynamic modelling

Jannis M. Hoch<sup>1,2</sup>, Jeffrey C. Neal<sup>3</sup>, Fedor Baart<sup>2</sup>, Rens van Beek<sup>1</sup>, Hessel C. Winsemius<sup>2,4</sup>, Paul D. Bates<sup>3</sup>, Marc F.P. Bierkens<sup>1,2</sup>

# GLOFRIM v2.0

- > Generic framework
- > Based on BMI + thin python layer for homogeneity
- > 2-step initialization
- > Support for several grid types
- > New models easily added (if BMI API), now supporting

>> PCRGLOB-WB, wflow (hydrological)

>> CaMa-Flood, Lisflood-FP, Delft3D-FM (hydrodynamic)

### GLOFRIM v2.0

#### **GLOFRIM configuration file example**

[models]
# reference to model config files
PCR=path/to/prcglobwb.ini
CMF=path/to/camaflood.nam

[coupling]
# time step for exchanges [sec]
dt=86400

[exchanges]
# PCR runoff [m] to CMF runoff [m]
PCR.runoff=CMF.roffin

#### **GLOFRIM runner script**

python glofrim\_runner.py run /path/to/glofrim.ini

Hydrology -> Routing

## GLOFRIM v2.0

```
from glofrim import Glofrim
 1
 2
     from datetime import datetime
 3
     # initialize combined bmi
 4
     cbmi = Glofrim()
 5
 6
 7
     # initialize configuration
 8
     cbmi.initialize config(config fn='path/to/config.ini')
 9
     # possible to edit configuration. e.g. set start times accross models
10
     cbmi.set start time(datetime(2000,1,1))
11
12
     # initialize models
13
14
     cbmi.initialize model()
15
     # interact with individual models
16
17
     cbmi.bmimodels['PCR'].get value('variable name')
18
19
     # run combined models & exchange data at each timestep
20
     cbmi.update until(cbmi.get end time())
21
```

### Next steps

> Release v2.0 with accompanying paper late-2018/ early 2019
> v2.x

>> Xarray based NetCDF adaptor. For 1 way coupling & to tap into online data!

>> Test two-way coupling schemes

>> Add regridding options (xesmf ?)

IVM Institute for Environmental Studies



# Questions ?? / Suggestions !!

#### GitHub



https://github.com/openearth/glofrim



https://glofrim.readthedocs.io/en/latest/

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