

HyMUSE: a multi-model framework for Hydrology

Inti Pelupessy

EGU General Assembly, Vienna, 08/04/2019



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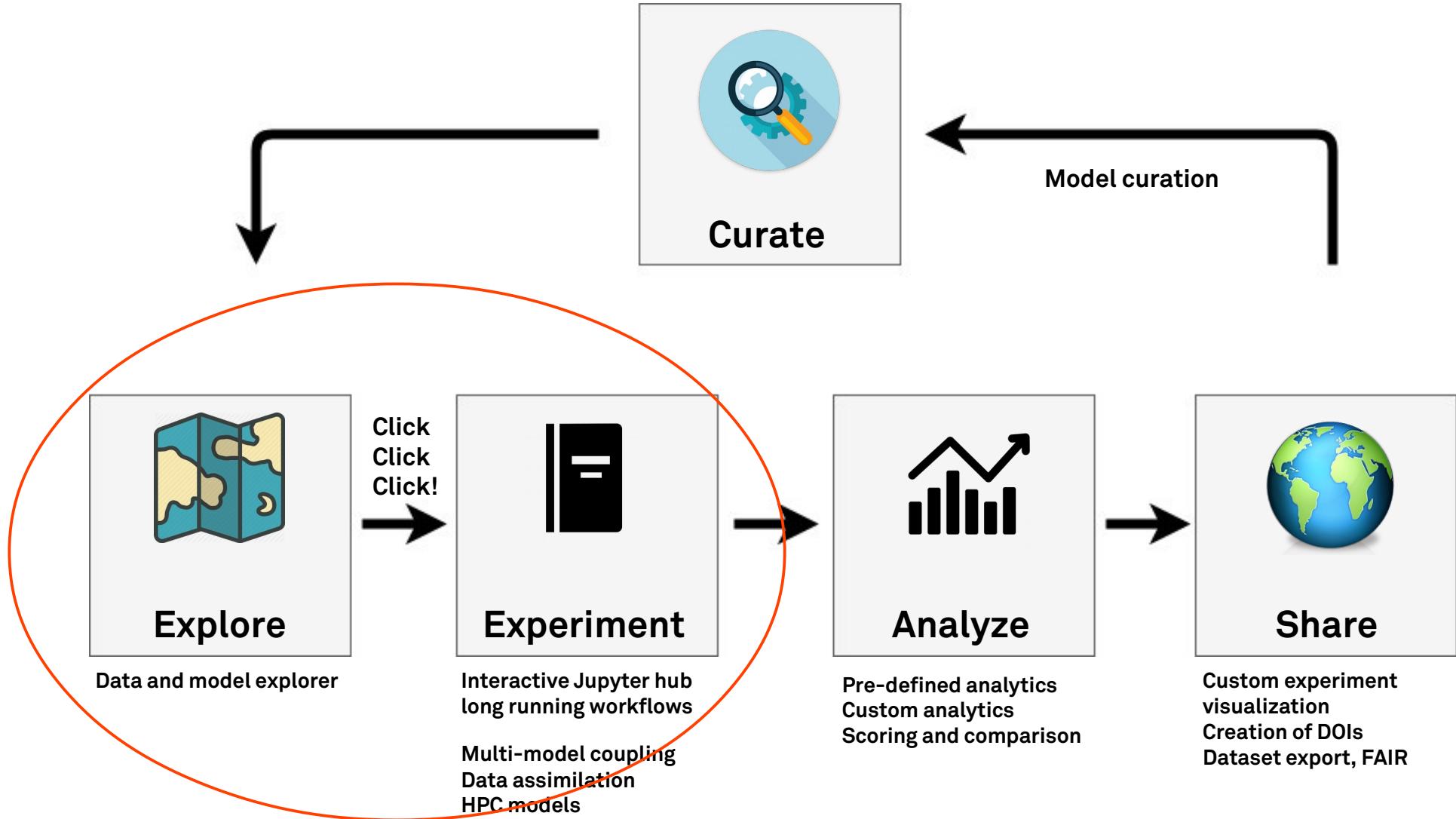
+ partners at Deltares,
Utrecht University,
Wageningen a.o.

eWaterCycle II Goals

The eWaterCycle II project aims to be a global-scale hydrological modeling platform that allows users to apply hydrological models to answer their research questions.

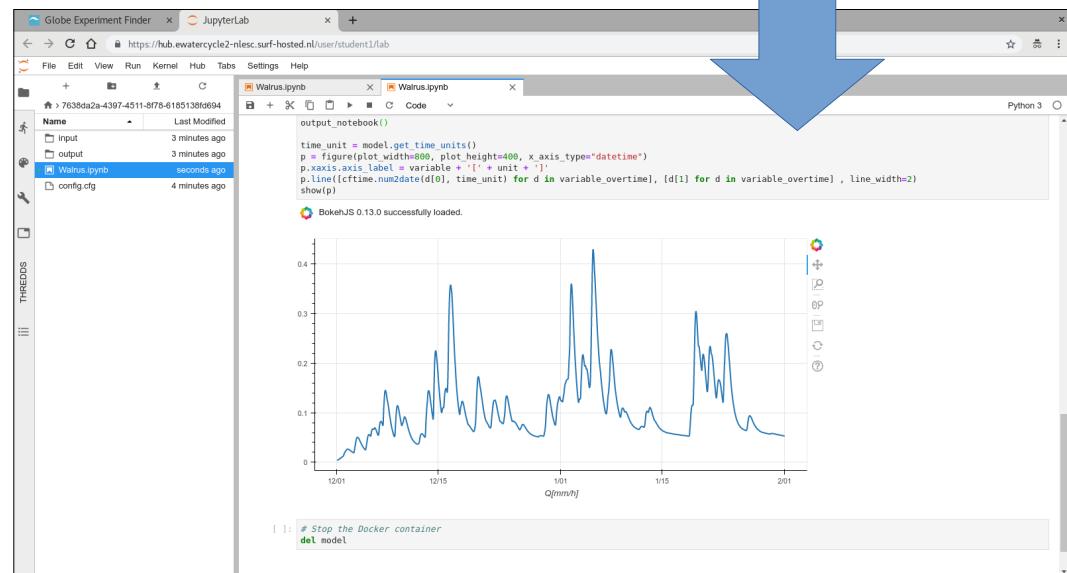
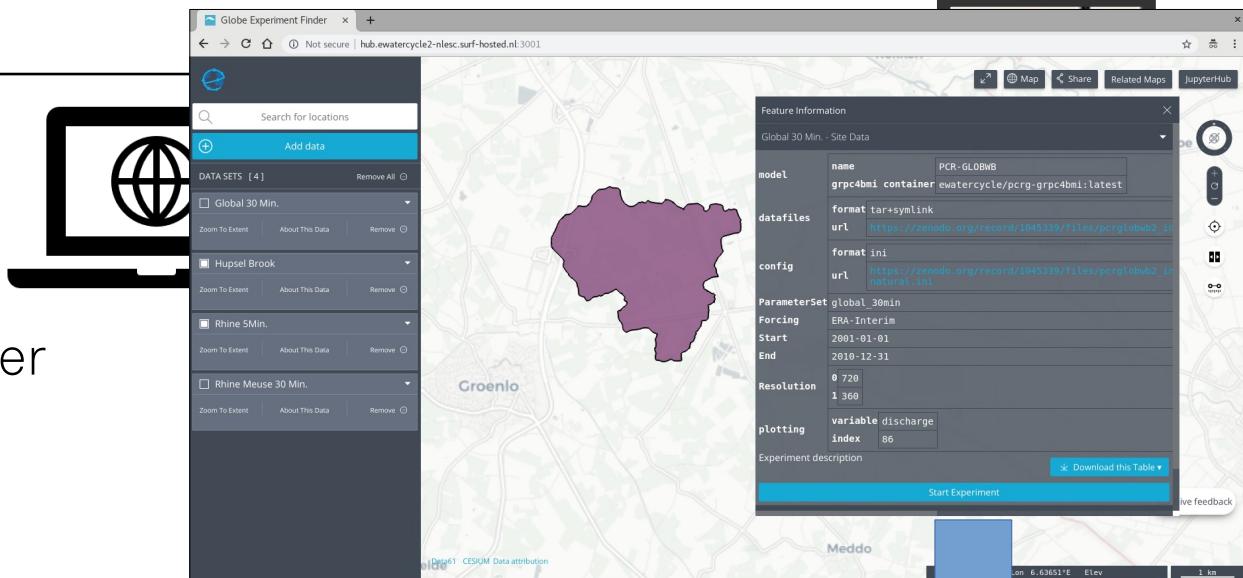
- Combine global hydrological models with local hydrological models, written in different programming languages,
- Recognize findable, accessible, interoperable, and reproducible (FAIR) practices.
- Make it much easier to work together on models.
- Cut down on fte's and time needed to run experiments.
- For this we need a lot of models and a lot of hydrologists to join in.

“Towards FAIR & Open Science in Hydrological Modelling”

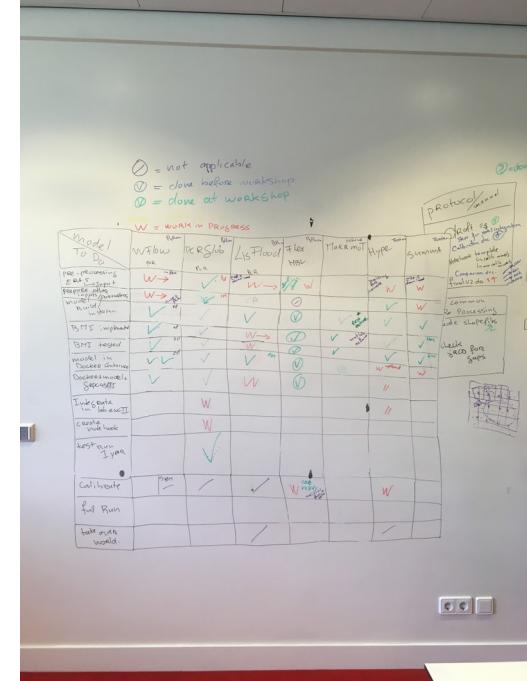


eWaterCycle II workflow

1. Choose a region in online viewer
2. Get list of available models / forcings / observations for that region.
3. Make a selection and “launch experiment”.
4. Get an (online!) Python notebook with the hello-world for that model: a hydrograph at the basin outlet compared to observations



Building a community



Lorentz workshop in April 2019 to incorporate seven hydrological models

introducing HyMUSE

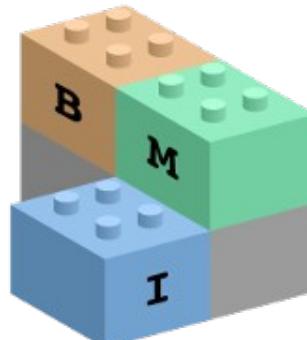
Problem: we need a method to interact with simulation codes that is:

- simple & intuitive to use
- easy to install
- interfaces codes written in different languages
- uses homogeneous interfaces
- allows the use non-local compute resources

Solution:



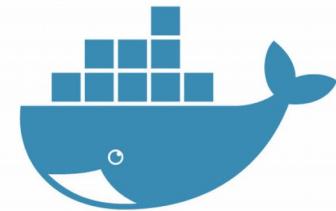
AMUSE/ OMUSE



Basic Model Interface



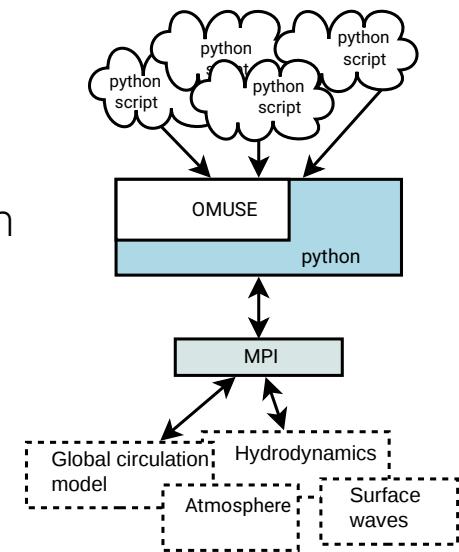
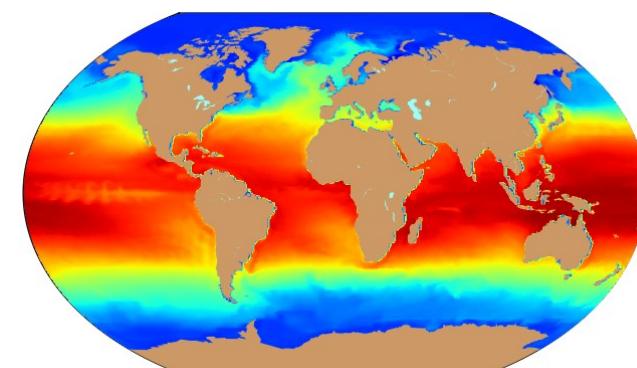
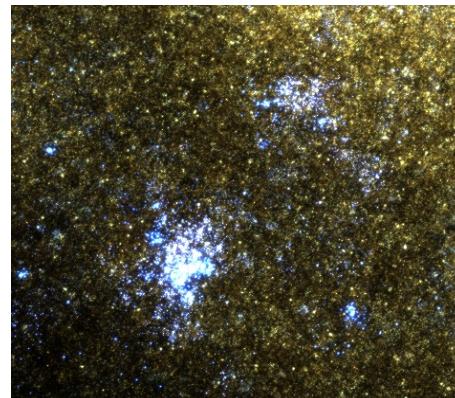
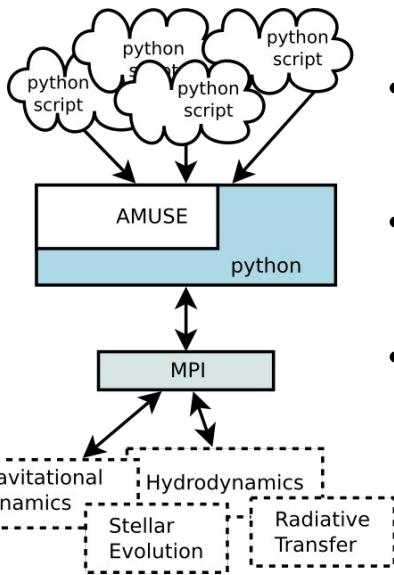
GRPC4BMI



containers

AMUSE & OMUSE

- model coupling frameworks for astrophysics & climate sciences (developed at Leiden Observatory, Utrecht University and the Netherlands eScience Center):
- provide a homogeneous Python environment to run community codes
- enable new code couplings and interactions between components
- facilitate multi-physics and multi-scale simulations



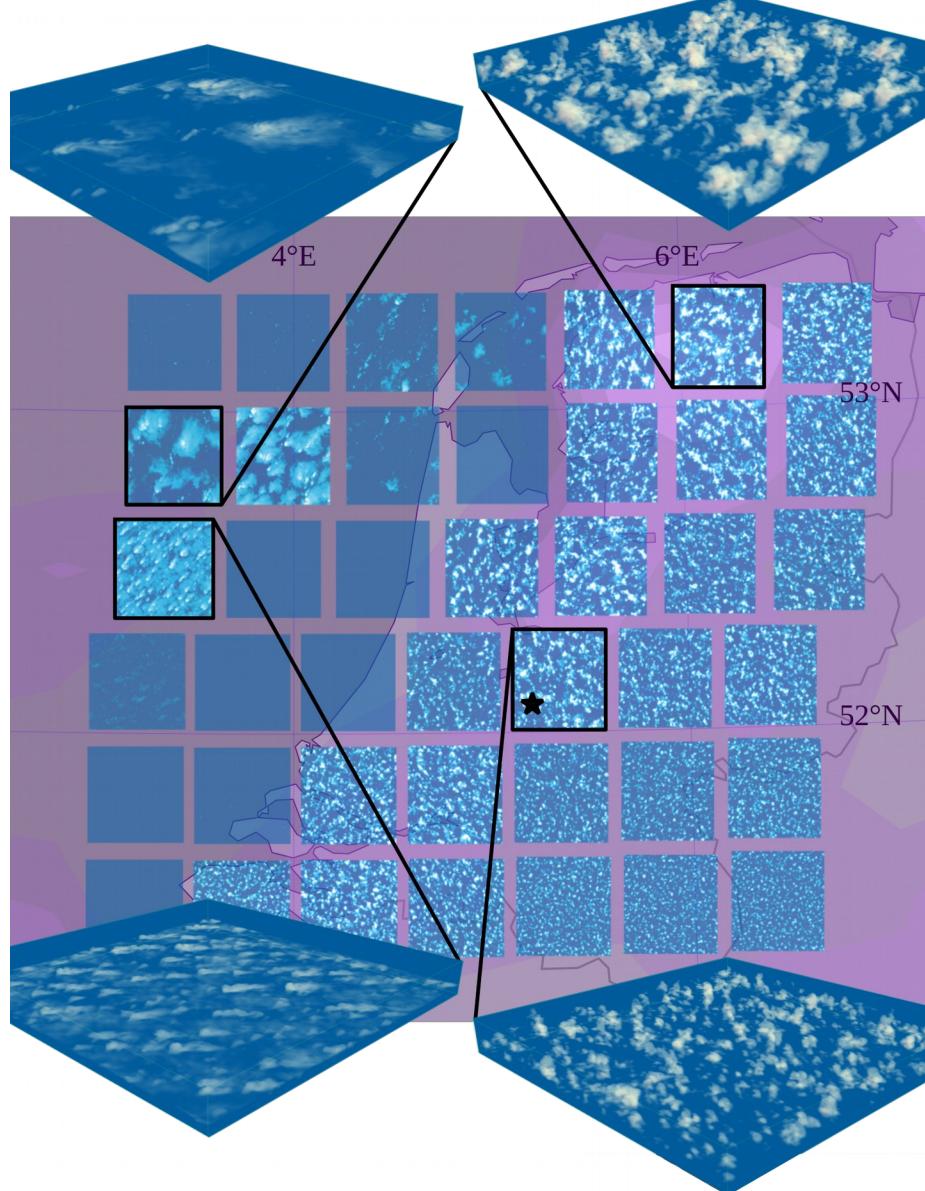
Pelupessy+ 2017
Pelupessy+ 2013

Example: DALES – OIFS coupling

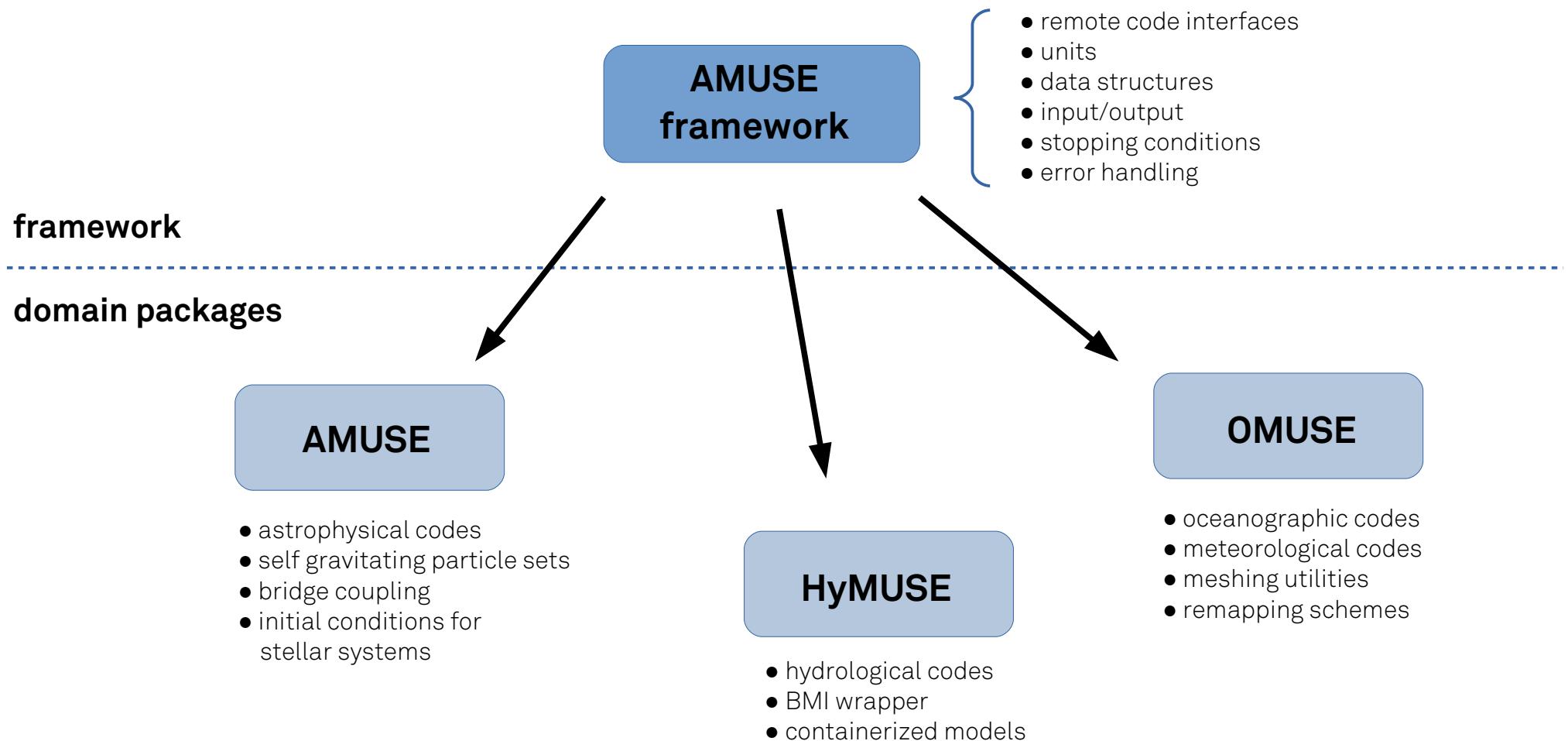
- superparametrization
- embedding of cloud resolving in global atmosphere model
- two way coupling
- N models in single global instance

see Jansson et al. #11303 (AS1.1)...

(and other examples:
Pelupessy et al. 2017, GMD 10, 3167)



ABC-MUSE architecture



'Hello Hydro-world'

```
“imports”      from hymuse.units import units
                from hymuse.community.pcrglobwb.interface import PCRGlobWB

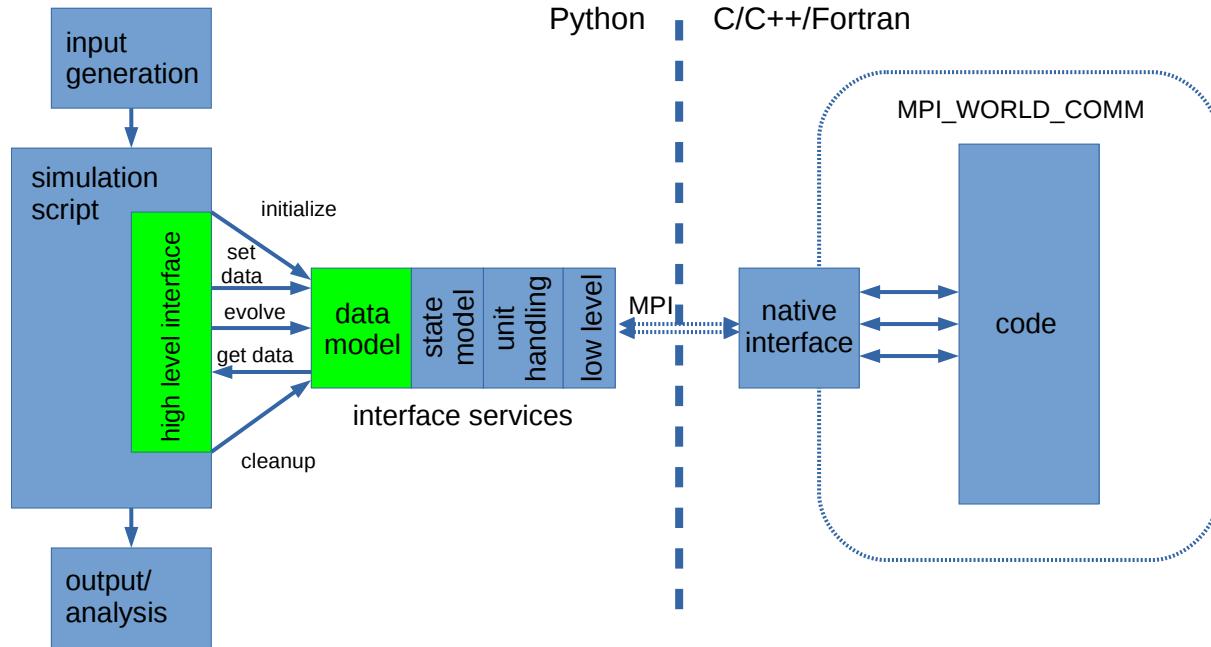
“instantiate code”    code=PCRGlobWB()

“initialize model”   code.parameters.ini_file="setup.ini"

“evolve”         code.evolve_model(code.model_time + (1.| units.day) )

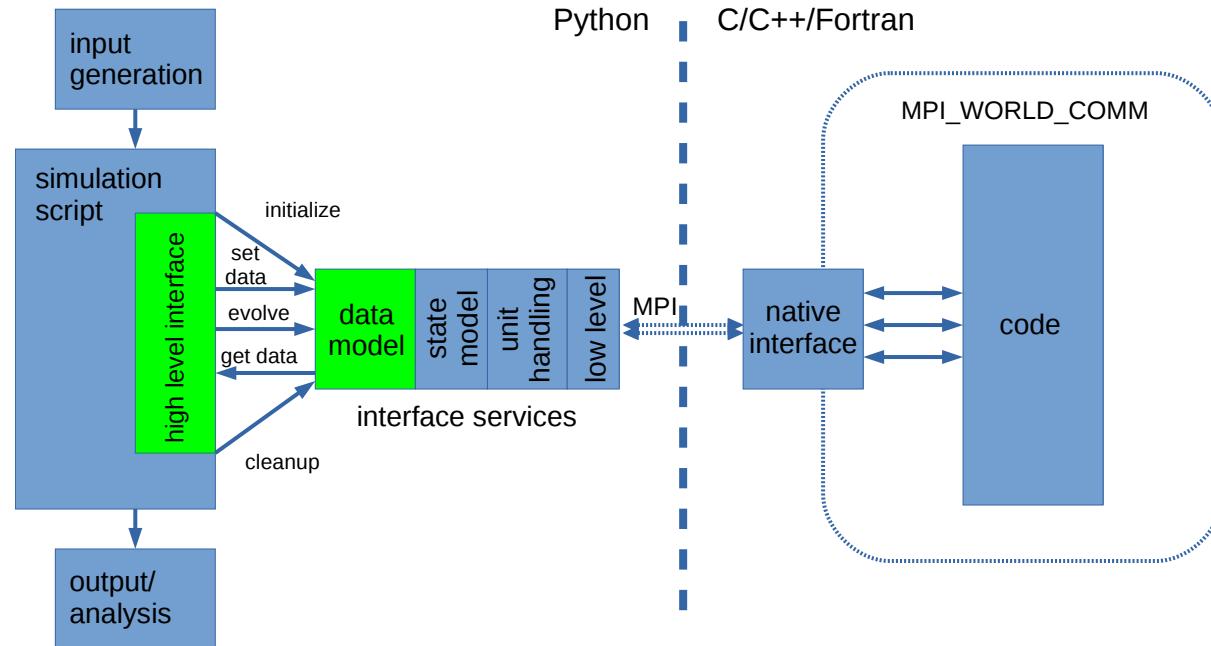
“analysis”       station=code.grid_0.samplePoint( lon= 6. | units.deg,
                                                lat= 51. | units.deg )
                print("discharge:", station.discharge.in_(units.m**3/units.s) )
```

The HyMUSE interface: a bird's eye view



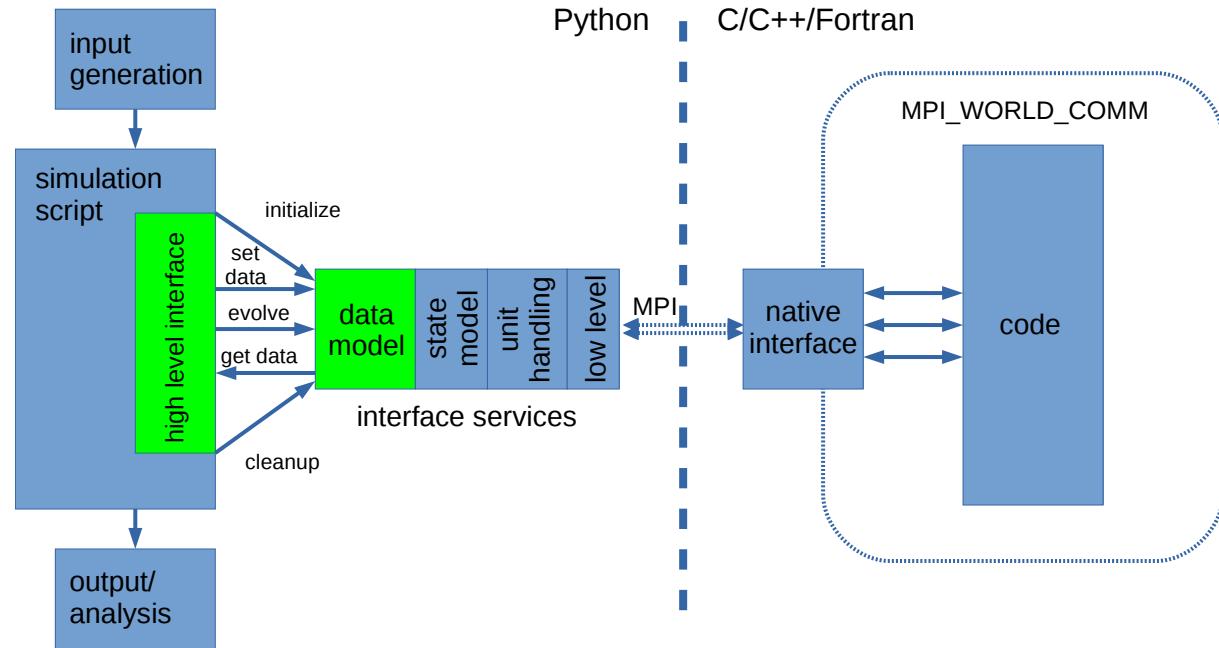
(i) HyMUSE is based on remote code interfaces

The HyMUSE interface: a bird's eye view



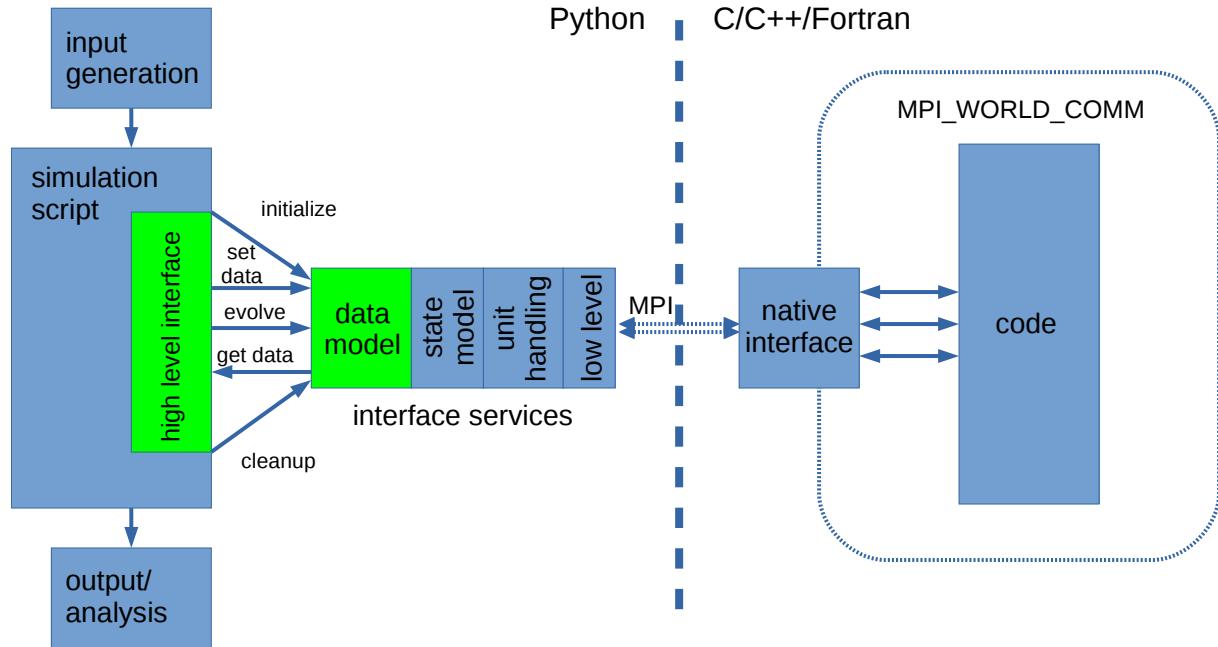
(ii) codes run in parallel & calls can be asynchronous

The HyMUSE interface: a bird's eye view



(iii) HyMUSE provides a number of *interface services*: unit handling, data structures, code state handling etc..

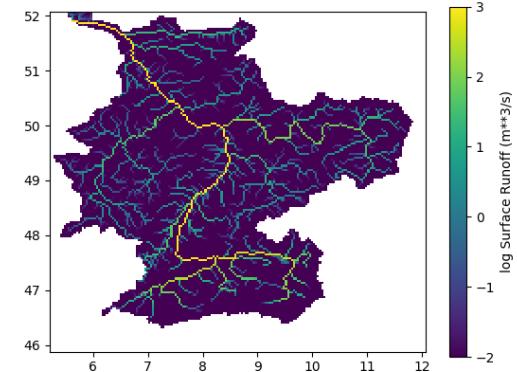
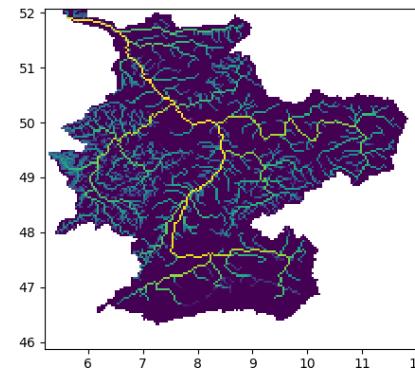
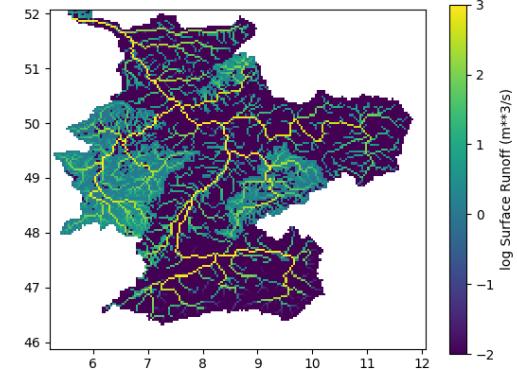
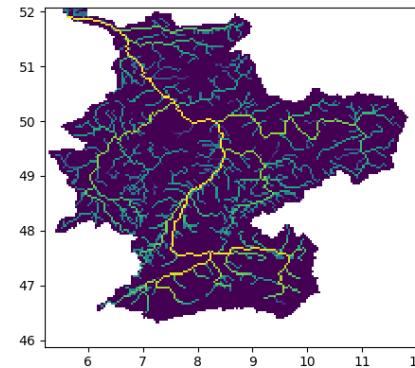
The HyMUSE interface: a bird's eye view



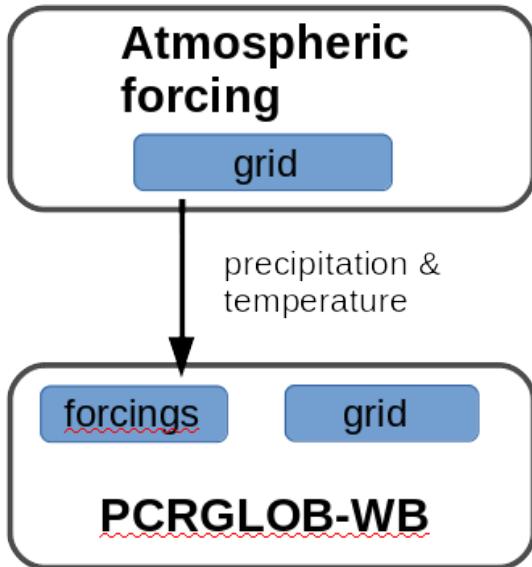
(iv) HyMUSE contributed codes can be containerized and based on BMI, allowing for easy interoperability with e.g. pyMT

What can you do with HyMUSE?

- simplify model setup and runs,
- scripting:
 - event detection,
 - stopping conditions
- 'online' data analysis
- ensemble simulations:
 - parameters searches
 - optimizations (e.g. MCMC)
 - data assimilation
- model comparison: running problems with different codes and methods
- coupling different codes to construct new solvers

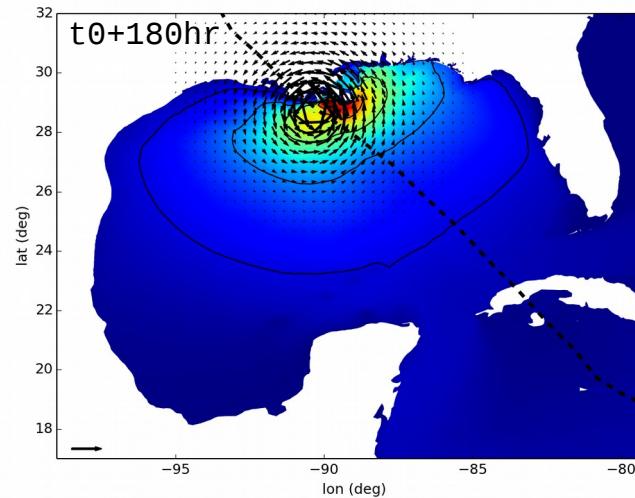
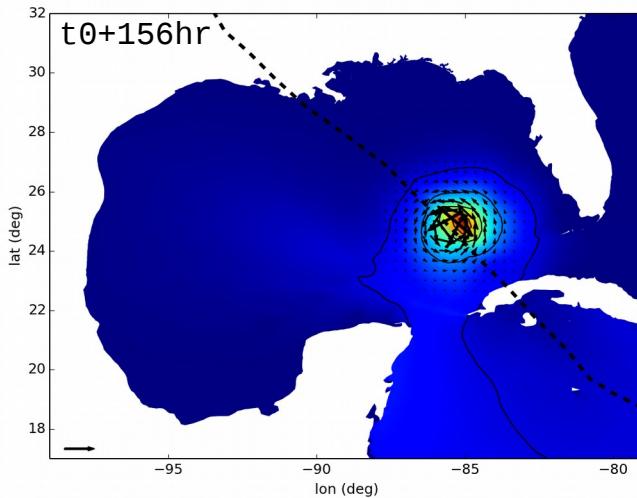


Code couplings with HyMUSE

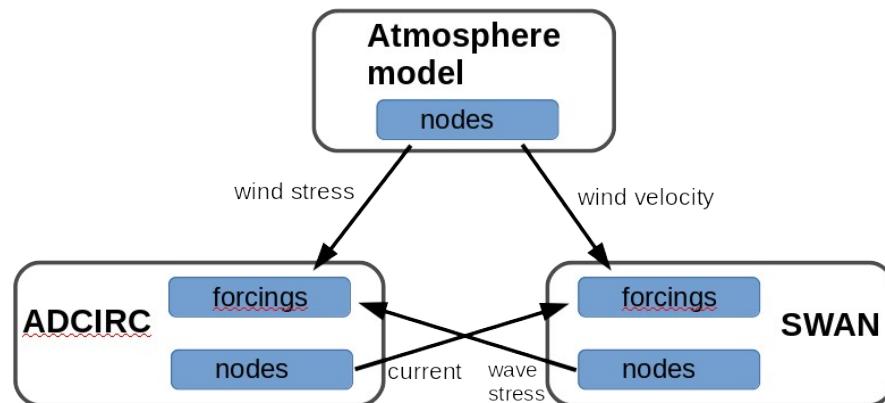


```
pcrglob = PCRGlobWB()  
  
p.parameters.ini_file = ini_file  
pcrglob.parameters.use_interface_forcings = True  
  
meteo = netcdf_meteo( "precipitation_2001to2010.nc",  
                      "temperature_2001to2010.nc" )  
  
channel = meteo.grid.new_channel_to( pcrglob.forcings )  
  
while pcrglob.model_time<tend:  
    meteo.evolve_model( pcrglob.model_time + dt )  
    channel.copy_attributes( ["precipitation", "temperature"] )  
    pcrglob.evolve_model( pcrglob.model_time + dt )
```

Hurricane Gustav with a coupled hydrodynamic/ wave model



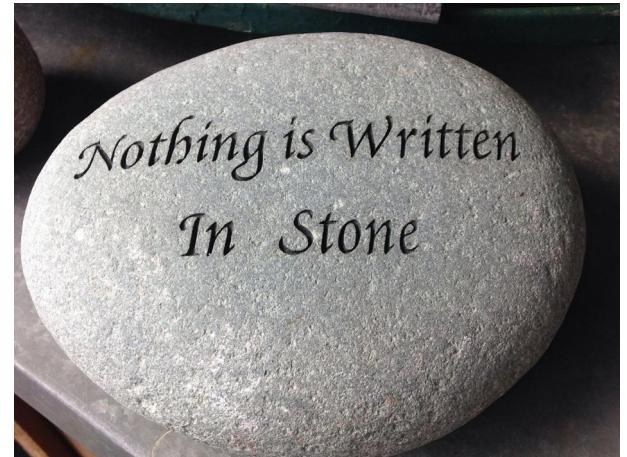
(Pelupessy+ 2017)



```
channel1=hurricane.grid.new_channel_to( swan.forcings )
channel2=hurricane.grid.new_channel_to( adcirc.forcings )
channel3=adcirc.nodes.new_channel_to( swan.forcings )
channel4=swan.nodes.new_channel_to( adcirc.forcings )
while time<tend:
    hurricane.evolve_model(time+dt/2)
    channel1.copy_attributes(["vx", "vy"])
    channel2.copy_attributes(["tau_x", "tau_y"])
    adcirc.evolve_model(time+dt/2)
    swan.evolve_model(time+dt/2)
    channel3.copy_attributes(["current_vx", "current_vy"])
    channel4.copy_attributes(["wave_tau_x", "wave_tau_y"])
    time+=dt
```

current status

- basic capability
- refinement of high level interfaces needed
- extending community code base
- requirements for "eWaterCycle approved" codes?
- initialization, restarts
- development of BMI:
 - * match with eWaterCycle goals?
 - * Technical factors: limitations/ shortcomings
 - * Human factors: different standards floating around, different opinions on usage



Conclusions



- eWaterCycle in development as a FAIR platform for hydrology
- HyMUSE aims to be the flexible multi-model framework for the computational core
- It is in development at the Netherlands eScience Center
- Community driven: we need your input!

www.ewatercycle.org

lab.ewatercycle.org

github.com/ewatercycle/hymuse

Thank you!