

# Tightly-coupled workflows using **MUSCLE2**



22/11/2013

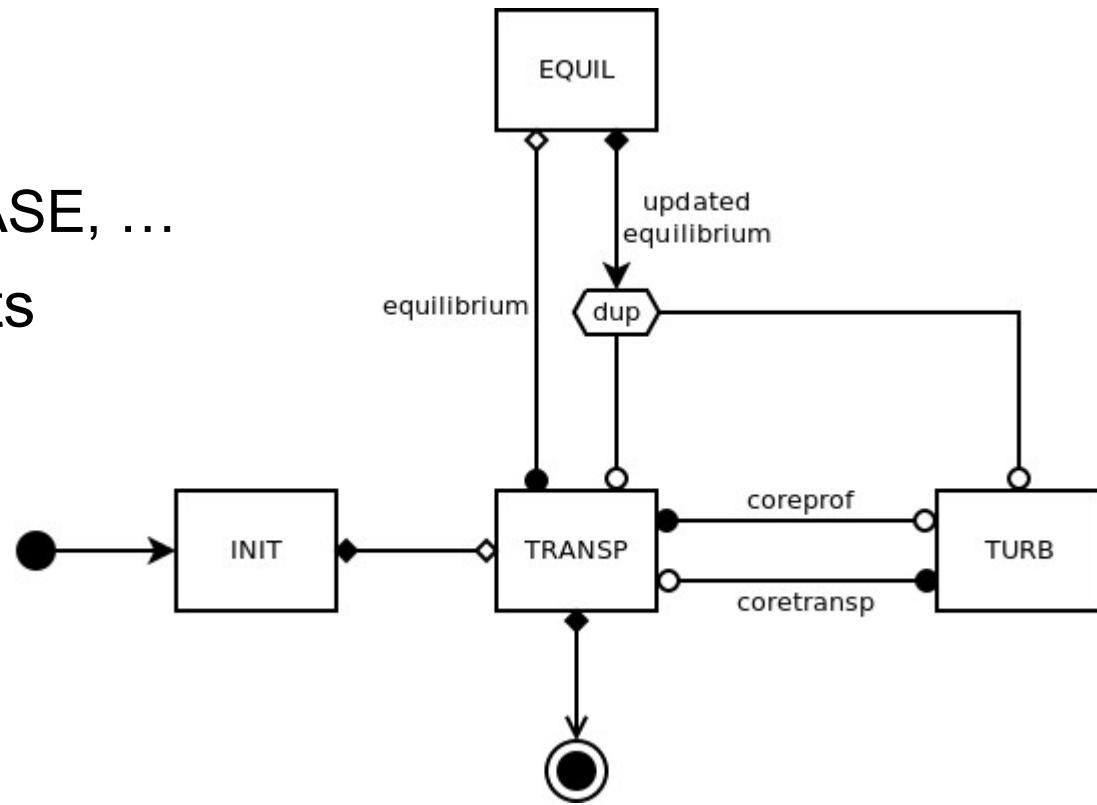
Olivier Hoenen

# Use case

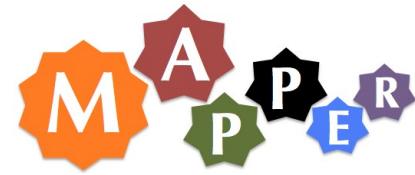


- Tightly-coupled (cyclic)

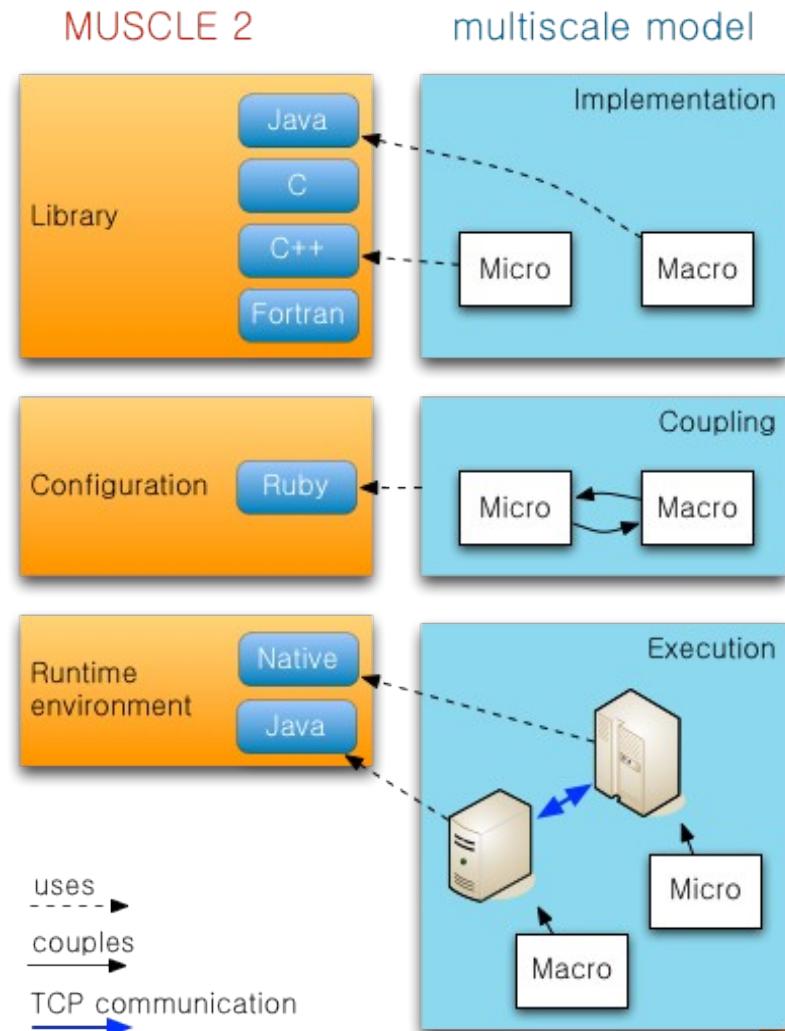
- Transport solver  
ETS
- Equilibrium code  
Helena, Bdseq, CHEASE, ...
- Transport coefficients  
Bohmgb, GEM, ...



# MUSCLE 2



- Coupling library
  - Core in Java (now also in C++)
  - API in C/C++, F90 (Python, Scala)
  - Based on “kernels”
- Handle time evolution
  - Reflects MML terminology
- Workflow described in Ruby (script)
- Inter-sites communications
  - MTO (TCP, MPWide)  
<http://apps.man.poznan.pl/trac/muscle>

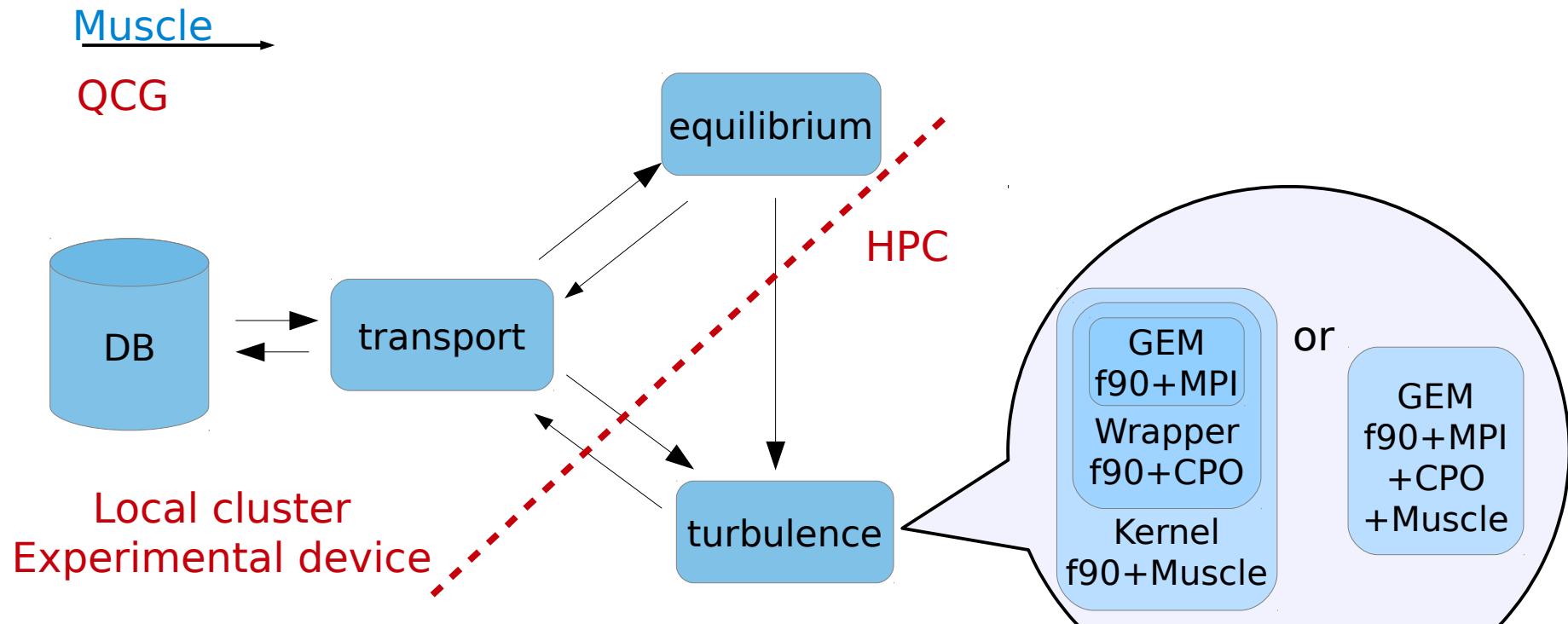


# Kernels



- API familiar to MPI users
  - `MUSCLE_Init`, `MUSCLE_Receive`,  
`MUSCLE_Get_Property`, `MUSCLE_Send`,  
`MUSCLE_Finalize`
- Receive/send handle accept basic C types
- Kernel contains time loop
  - `MUSCLE_Will_Stop` or hand-made
- Native source code → executable → **concurrent exec**
  - Scheduling is controlled by dataflow
- MPI kernels → only rank 0 calls MUSCLE

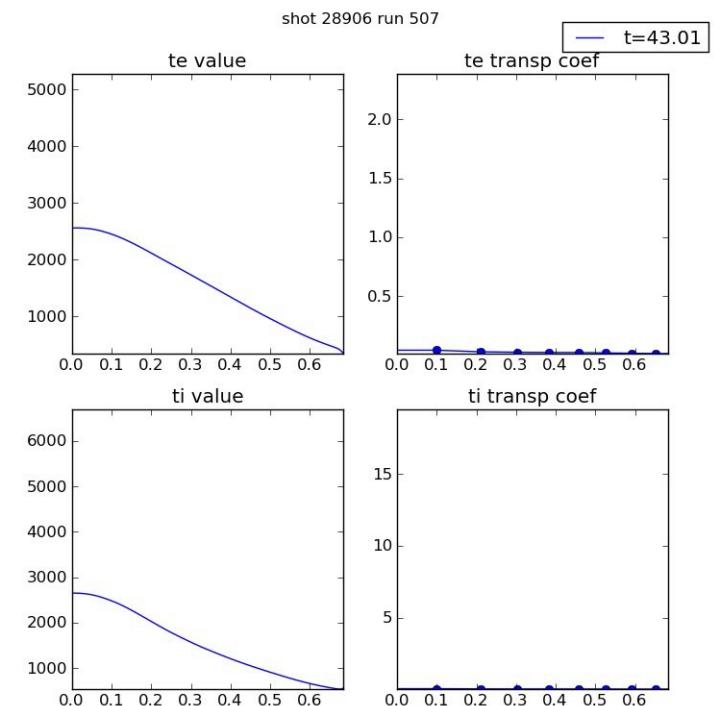
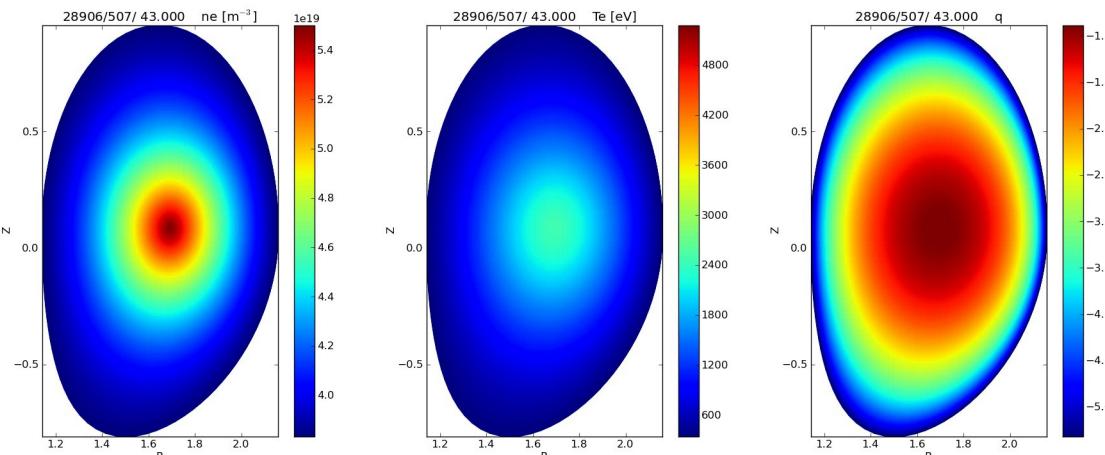
# Distributed run



# Some results

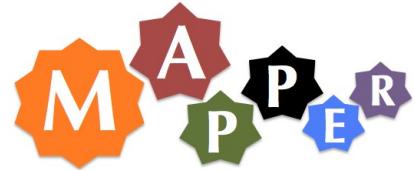


- Example on AUG shots:
  - 1s, transport dt=0.01, gyrofluid on 8 fluxtubes
  - 128 cores → wallclock = 10:36:26 (Gateway)





Questions?



# Ruby CXA file

- Describes the workflow
- Kernels id and path
  - User defined: Native, MPI, Java
  - Utils: duplication mapper,...
- Variables
  - Global or local to a kernel
- Links between kernels
  - Attach kernels, attach ports
  - Add filters: conversion, compression, etc...

```
# configuration of commands
cxa.env[ "transp:command" ]="ets_kernelB"
cxa.env[ "turb:command" ]="gem_kernelB"

# global params
cxa.env[ "time_step" ] = 0.1
# kernel specific params
cxa.env[ "init:cpo_file" ] = "CPO_000004_000001"
cxa.env[ "transp:solver_type" ] = 3
cxa.env[ "transp:sigma_source" ] = 0

# declare kernels
cxa.add_kernel('transp',
'muscle.core.standalone.NativeKernel')
cxa.add_kernel('turb',
'muscle.core.standalone.MPIKernel')
cxa.add_kernel('dupEquil',
'muscle.core.kernel.DuplicationMapper')

cs.attach('transp' => 'turb') {
tie('coreprof_out','coreprof_in')
}
cs.attach('turb' => 'transp') {
tie('coretransp_out','coretransp_in')
}
```

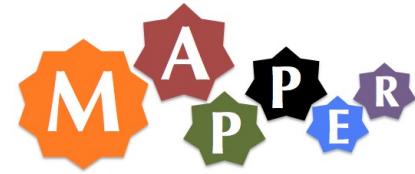
# QCG-Broker



- Resource management and brokering service
  - Assign kernels to site
  - Processes counts
  - Advance reservation
- Additional scripts
  - Preprocessing
  - Postprocessing
- Environment
  - stderr/stdout
  - modules

```
<task persistent="true" taskId="task">
  <requirements>
    <topology>
      <processes masterGroup="true" processesId="transp">
        <processesCount> ...
        <candidateHosts>
          <hostName>inula.man.poznan.pl</hostName>
        </candidateHosts>
      </processes>
      <processes processesId="turb">
        <processesCount> ...
        <candidateHosts>
          <hostName>zeus.cyfronet.pl</hostName>
        </candidateHosts>
      </processes>
    </topology>
  </requirements>
  <execution type="mapper">
    <executable>
      <application name="muscle2"/>
    </executable>
    <arguments>
      <value>ParallelModelsB.cxa.rb</value>
    </arguments>
    <stdout> ...
    <stderr> ...
    <stageInOut> ...
    <environment> ...
  </execution>
  <executionTime> ...
</task>
```

# On Helios



- First benchmark on Helios:
  - Scalability follows main component performance (GEM)
  - Speed-up loss from 512 to 1024 due to impact of serial components : ETS+CHEASE takes around 10-15s
  - At 2048 cores and above, non scalable behavior is due to non-scalable I/O usage (naïve implementation of serialization, etc...): to be investigated

