

Open MOdeL Experiment

A generic workflow platform for distributed experiments on complex-system models

__arge scale computing

- Compute transparently on multi-core machines, clusters, and grids: no particular knowledge is required to access massively parallel environments
- Embed your own programs in less than 5 clicks (C, C++, Java, Scala, Scilab, R, Python, Netlogo, ...)
- Design your distributed process using a powerful workflow approach (loops, alternative processing, massive parallelism)
- Prototype your workflow locally and take advantage of the parallelism with no extra effort: the Zero-deployment approach does not require any installation step on remote machines
- Use OpenMOLE as a scalable computing engine for your web portal (ex: bioemergences.eu is powered by OpenMOLE)



- Embed your model
- Design your model exploration:
 - > design of experiments,
 - > stochastic model replication,
 - > calibration,
 - > sensitivity analysis,
 - > ...
- Distribute the numerous executions of your model to explore it:
- > on your multi-core computer // speed up around 4
- > on a multi-core server // speed up around 50
- on a cluster // speed up arround 200
- > on a grid // speed up > 1000
- Reuse your experiments on several models





Subscribe to the newsletter : users@list.openmole.org

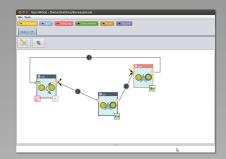


OpenMOLE: for whom?

• Are you a computer scientist?
Script your workflow!

```
val i1 = Prototype[Int]("i1")
val i2 = Prototype[Int]("i2")
val j = Prototype[Int]("j")
val hello = GroovyTask("hello", "j = Model.compute(i1, i2)")
hello addInput i1
hello addInput i2
hello addUnput j
hello addUnput j
hello addUnput j
hello addUnput i2
hello addUnput j
hello addUnput i2
hello addUn
```

Do you make typos, or do you prefer graphical applications?
Use the click and run Application!



•Do you need large scale computing facilities for your web site Use the OpenMOLE web-server!

