

Data Analysis Sequencing - EDNA



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Why do we need EDNA?

- EDNA is the best answer we (developers) have come up with so far for answering these questions :
 - How can we automate data analysis workflows?
 - “pipeline” existing scientific software for (online) data analysis workflows
 - abstract certain calculations to be “generic”, e.g. indexing of a diffraction pattern
 - “flexible” workflows, rapid changes depending on the scientific needs
 - How can we make these workflows robust?
 - easily adapted to new versions of scientific software packages
 - How can we collaborate efficiently?
 - re-use of code without breaking existing functionality

What is EDNA?

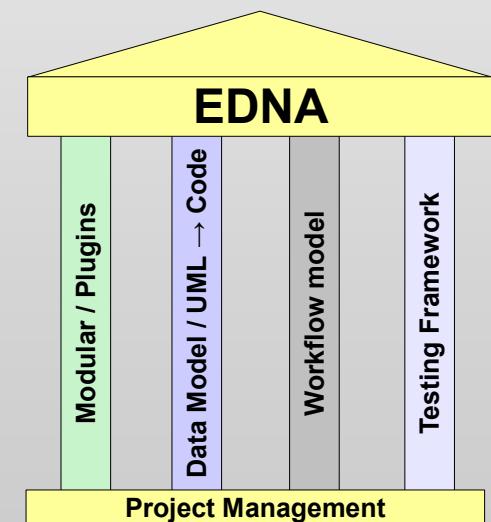
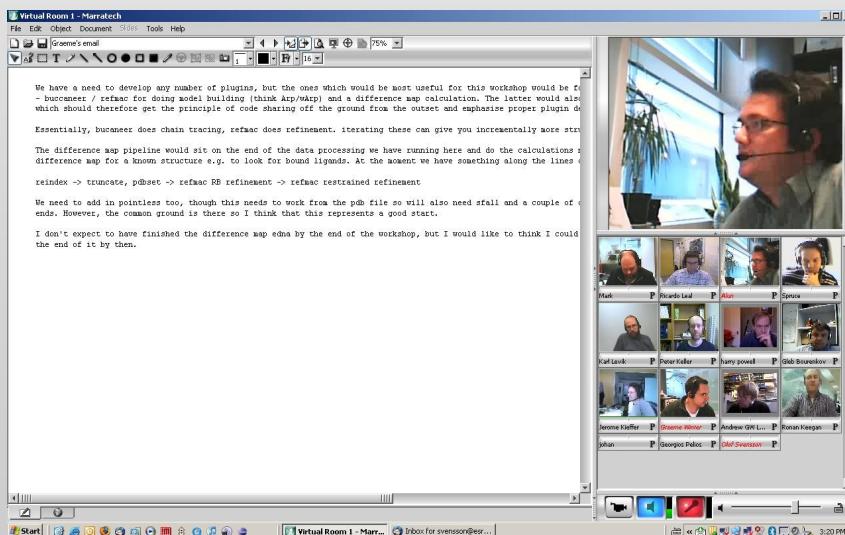


- EDNA is about collaboration:

- Code sharing (SVN)
- Coding conventions
- Code reviews
- Open source (LGPL, GPL)
- Bug tracker
- Wiki : <http://www.edna-site.org>
- Memorandum of Understanding
- Executive committee
- Project manager / coordinator
- Regular meetings / video conferences

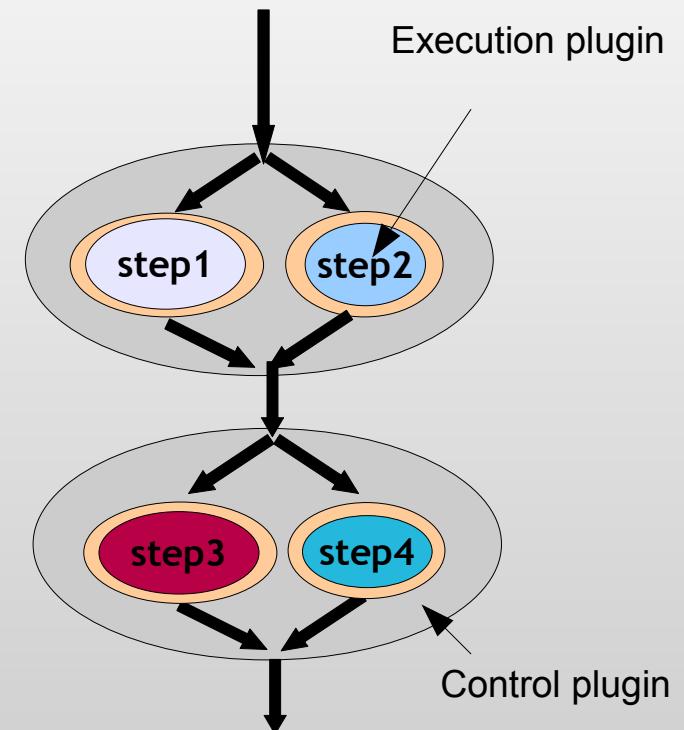
- EDNA is a framework:

- “Generic” kernel
 - Data modelling framework
 - Support for multi-threaded modules (plugins) development
 - Support for workflow development
 - Testing framework
- “Specific” applications (MXv1, bioSaxs etc.)
- Automatic testing and nightly builds
- Automatic API doc generation
- No GUI



EDNA Modularity : Plugins and their hierarchy

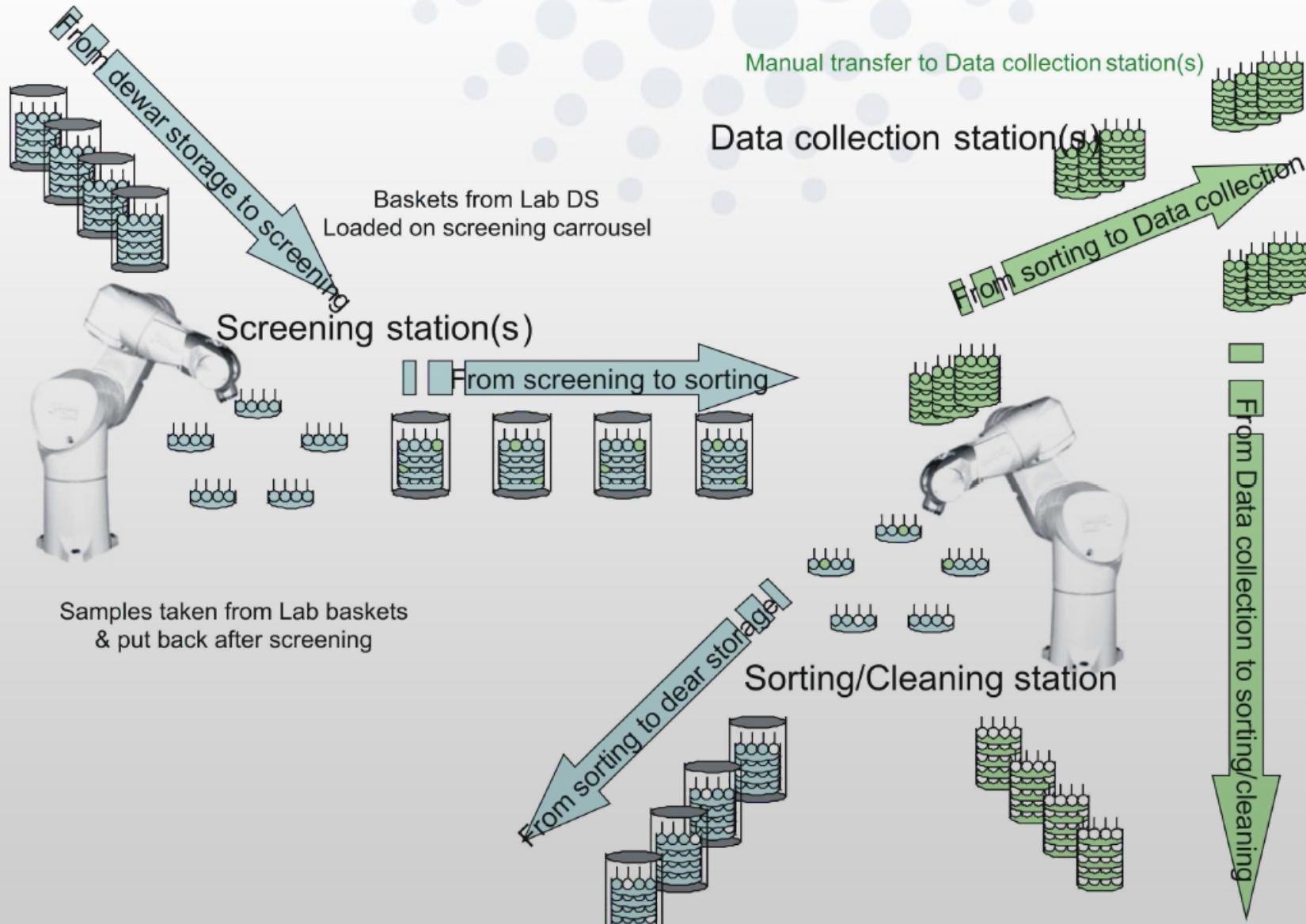
- Plugin base class :
 - Configuration, working directory, etc.
- Execution plugins :
 - Execution of external programs, e.g. (bash) scripts
- Controller plugins:
 - Control of execution plugins
 - Parallel execution
 - Synchronisation



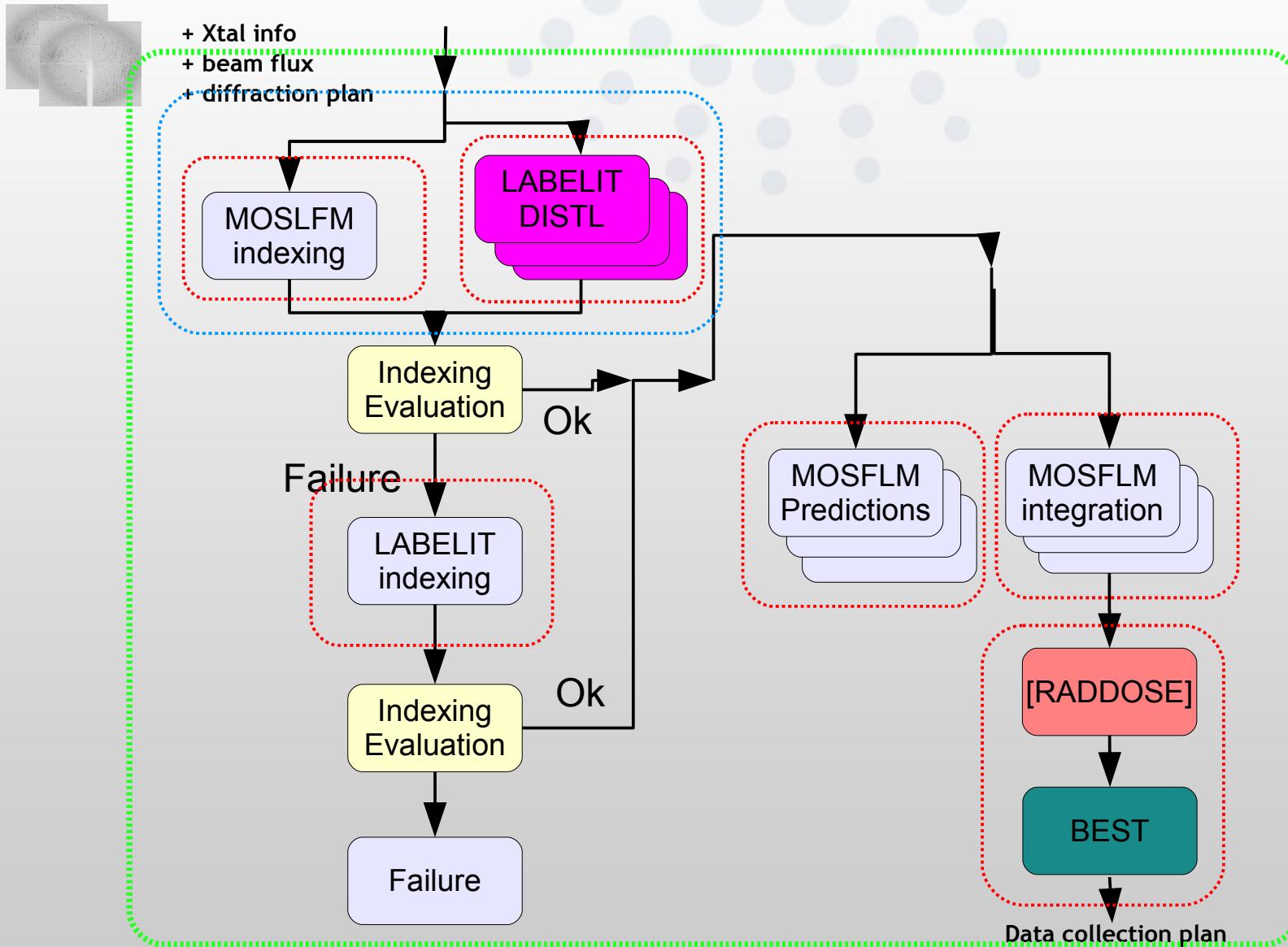
Existing scientific EDNA workflows

- Macromolecular crystallography:
 - Characterisation taking into account radiation damage (MOSFLM, Labelit, RADDOSE, BEST)
 - Connection with experiment data base (ISPyB)
 - Parallel execution of characterisation (GRID data processing)
 - Parallel creation of image thumbnails
- Diffraction Computed Tomography
 - SPD: Image correction, fast azimuthal integration
 - Sinograms saved in HDF5 format
- Small Angle Scattering
 - Image correction and fast azimuthal integration
- Full Field XAS
 - Image correction (dark, flat)
 - Image alignment (offset measurements by FFT)
 - HDF5 output

Challenge for the ESRF Upgrade : Massively Automated Sample Selection Integrated Facility



MXv1 Characterisation v1.2



EDNA Testing Framework

```
[UnitTest]: #####
[UnitTest]: Result for EDTestSuiteKernel : SUCCESS
[UnitTest]:
[UnitTest]: Number of executed test suites in this test suite : 2
[UnitTest]:
[UnitTest]:
[UnitTest]: Total number of test cases executed with SUCCESS : 14
[UnitTest]: Total number of test cases executed with FAILURE : 0
[UnitTest]:
[UnitTest]: Total number of test methods executed with SUCCESS : 48
[UnitTest]: Total number of test methods executed with FAILURE : 0
[UnitTest]:
[UnitTest]: Runtime : 25.847 [s]
[UnitTest]: #####
```

EDNA Testing Framework

```
[UnitTest]: #####
[UnitTest]: Result for EDTestSuitePluginUnitAll : SUCCESS
[UnitTest]:
[UnitTest]: Number of executed test suites in this test suite : 5
[UnitTest]:
[UnitTest]:
[UnitTest]: Total number of test cases executed with SUCCESS : 83
[UnitTest]: Total number of test cases executed with FAILURE : 0
[UnitTest]:
[UnitTest]: Total number of test methods executed with SUCCESS : 202
[UnitTest]: Total number of test methods executed with FAILURE : 0
[UnitTest]:
[UnitTest]: Runtime : 122.892 [s]
[UnitTest]: #####
```

EDNA Testing Framework

```
[UnitTest]: #####
[UnitTest]: Result for EDTestSuitePluginExecuteAll : FAILURE
[UnitTest]:
[UnitTest]: Number of executed test suites in this test suite : 5
[UnitTest]:
[UnitTest]:
[UnitTest]: Total number of test cases executed with SUCCESS : 124
[UnitTest]: Total number of test cases executed with FAILURE : 8
[UnitTest]:
[UnitTest]:
[UnitTest]: OBS! The following test methods ended with failure:
[UnitTest]:
[UnitTest]: EDTestCasePluginExecuteControlCharForReorientationv2_0_noKAPPA_ :
[UnitTest]:   testExecute :
[UnitTest]:     Plugin failure assert: should be False, was True FAILURE: Expected different from obtained - identifier
/mntdirect/_scisoft/users/svensson/tmp/EDTestSuitePluginExecuteAll_20110114-093729/tmpPW7olu
[UnitTest]:
...
[UnitTest]:
[UnitTest]:
[UnitTest]: Total number of test methods executed with SUCCESS : 129
[UnitTest]: Total number of test methods executed with FAILURE : 8
[UnitTest]:
[UnitTest]:           Runtime : 1108.997 [s]
[UnitTest]: #####
```

EDNA Testing Framework

Checking out EDNA Repository... to revision 2758

Tests using the default python under linux: Python 2.6.5

Launching EDTestSuiteKernel with /usr/bin/pythonSUCCESS! 16.411 [s]

Making EDNA-kernel tarball distribution.....

Launching EDTestSuiteCCP4v0 with /usr/bin/pythonSUCCESS! 16.466 [s]

Making CCP4v0 tarball distribution.....

Launching EDTestSuitePluginExecPlugins with /usr/bin/pythonSUCCESS! 920.778 [s]

Making ExecPlugins tarball distribution.....

Launching EDTestSuiteBioSaxs with /usr/bin/pythonSUCCESS! 49.655 [s]

Making BioSaxsv1 tarball distribution.....

Launching EDTestSuitePluginUnitMXPluginExec with /usr/bin/pythonSUCCESS! 57.916 [s]

Launching EDTestSuitePluginExecuteMXPluginExec with /usr/bin/pythonSUCCESS! 365.850 [s]

Launching EDTestSuitePluginUnitMXv1 with /usr/bin/pythonSUCCESS! 73.398 [s]

Launching EDTestSuitePluginExecuteMXv1 with /usr/bin/pythonSUCCESS! 3040.207 [s]

Making MXv1 tarball distribution.....

Tests using jython under linux: Jython 2.5.2rc2

Launching EDTestSuiteKernel with /home/tester/bin/jythonSUCCESS! 80.371 [s]

Tests using the default python under : WinePython 2.7.1

Launching EDTestSuiteKernel with /home/tester/bin/pythonwSUCCESS! 44.725 [s]

How EDNA will evolve in the future

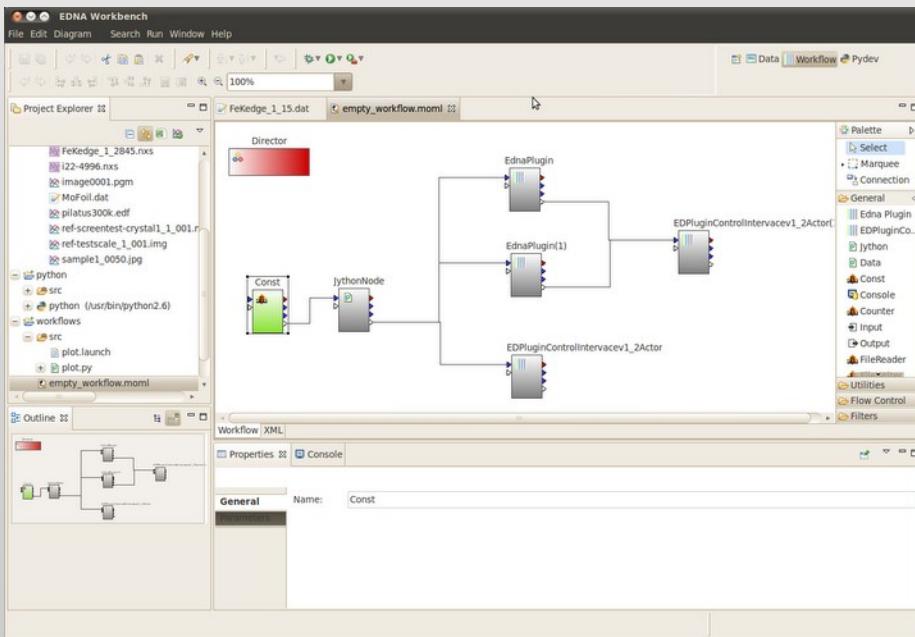
- Common EDNA developments (Kernel):
 - Improvements of the data model framework (in progress)
 - Improvements of logging (in progress)
 - Full support of Windows and MacOS (in progress)
 - Enhanced support of grid engines / job schedulers
 - Improved documentation (plugin use cases)
 - Graphical workflow editor (Data Analysis Workbench)
- Scientific developments:
 - MX further enhancements of characterisation (kappa, XDS etc)
 - MX auto processing wrappers
 - Biosaxs data analysis (EMBL Hamburg software suite)
 - Tomography
 - More to come...

How to run EDNA on a cluster with a job scheduler

- EDNA is optimised for parallel execution of processes/plugins:
 - Thread safe parallel execution of plugins
 - Automatic synchronisation
 - Automatic workload limitation
- Implemented : EDNA TANGO server
 - No job scheduler → no load balancing
- Today the EDNA kernel provides a limited support for running workflows on a cluster with a job scheduler:
 - Works only if the program is started by EDNA in a script
 - Call to grid engine submission must be blocking (sun/oracle grid engine “-sync y”)
- Future : support for any job scheduler (sun/oracle, torque, condor, pbs, oar etc)
 - Synchronisation through sockets?
 - Persistent EDNA plugin launchers?

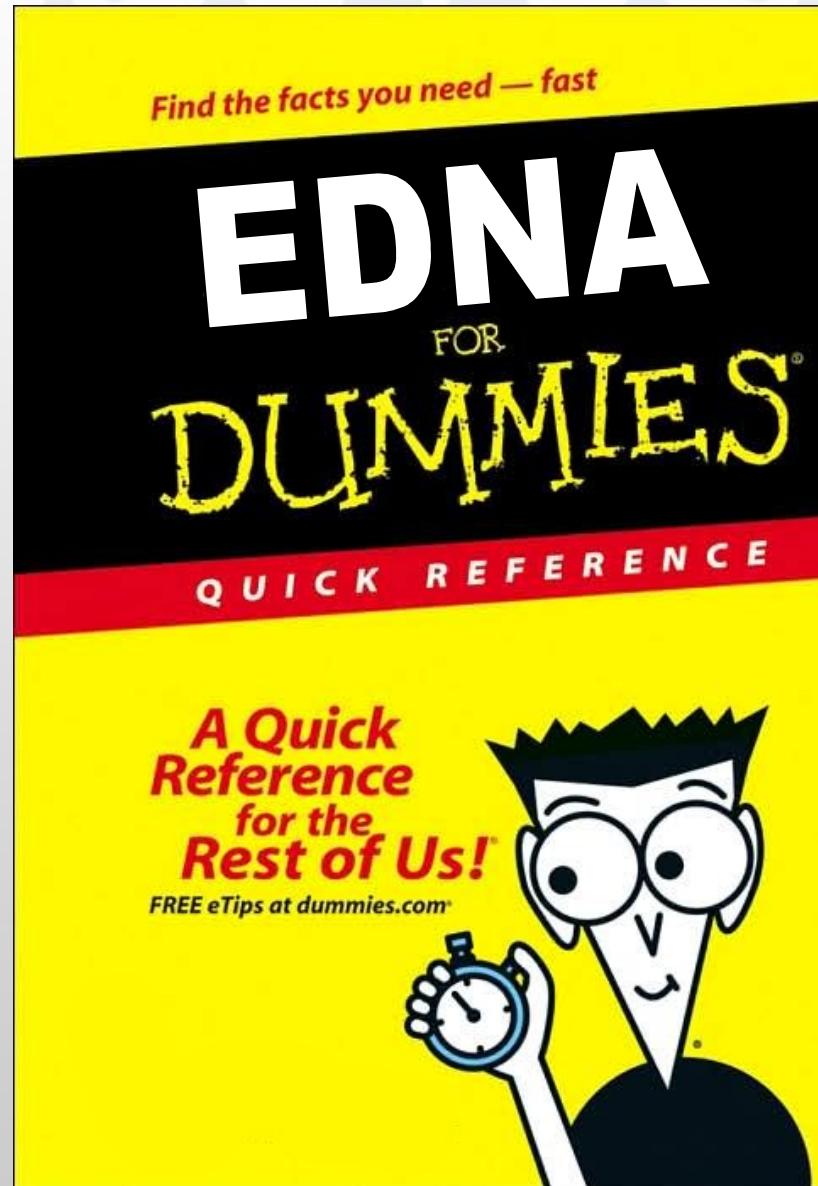
What the EDNA GUI could look like

- EDNA has no GUI framework
- All EDNA workflows must be executable on the command line
- Possible solution for a generic EDNA GUI:



- Workflow editor:
 - Implicit documentation of workflow
 - Implicit parallel workflows
 - Possibility to “easily” modify / construct new workflows
 - Possibility to debug workflows
 - Possibility to restart a stopped workflow

Documentation!



EDNA Documentation

- Available today :
 - Data models (png)
 - Automatic API doc generation
 - Wikipages with developers' "How-to's"
 - Minutes / presentations of previous meetings, code camps etc
- Planned :
 - Automatic plugin documentation repository (use cases etc)
 - Workflow documentation (workflow tool)

EDNA Collaborators 2010

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Alun Ashton^(b)

Andrew Leslie^(h)

Andrew McCarthy^(c)

Andrew Thompson^(k)

Clemens Schulze^(j)

Clemens Vonrhein^(f)

Darren Spruce^(e)

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(l) University of Sydney, Australia

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EDNA developers
Executive committee