



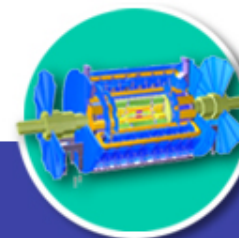
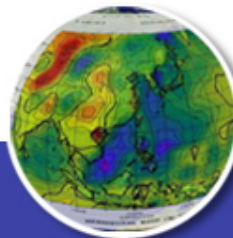
Ganga

The Job Submission Tool

WeiLong Ueng
wlueng@twgrid.org

www.euasiagrid.org

www.euasiagrid.eu





Objectives

- **This tutorial gives users to understand**
 - Why require Ganga in Grid environment
 - What advantages of Ganga
 - The Architecture of Ganga
 - The Implementation of Ganga
 - How to use Ganga for job management



Outline



- **Overview**
- **Ganga Job Components**
- **Ganga Architecture**
- **Ganga Interface**
- **Ganga & DIANE**
- **Ganga - Use Case**



Background

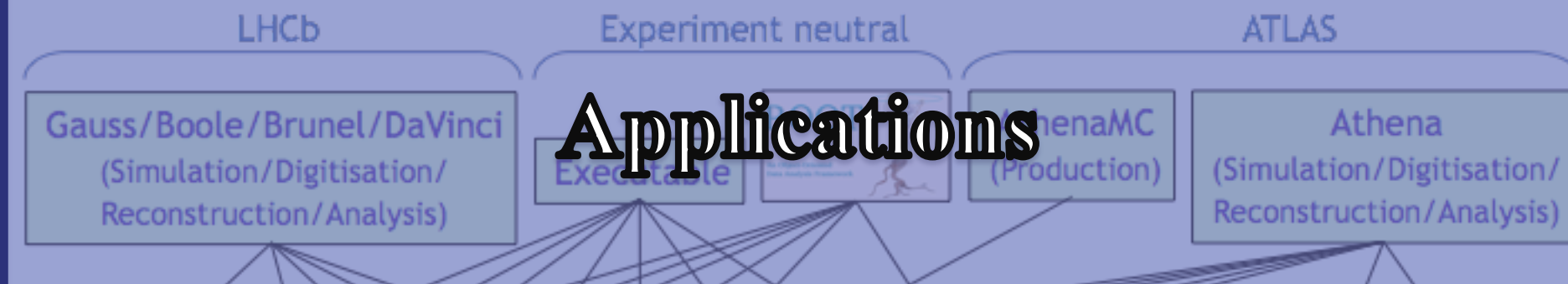


- **Growing numbers of distributed system**
 - Scaling of distributed systems, from local batch systems and community-specific services to generic, global Grid infrastructures
 - Debug on PC, then perform small-scale testing in local resources and finally run at full-scale in globally distributed Grids
- **Supports Different Backends**
 - Backends: PBS, Condor, LCG, EDS... etc
 - How to deploy on various backends in an efficient way

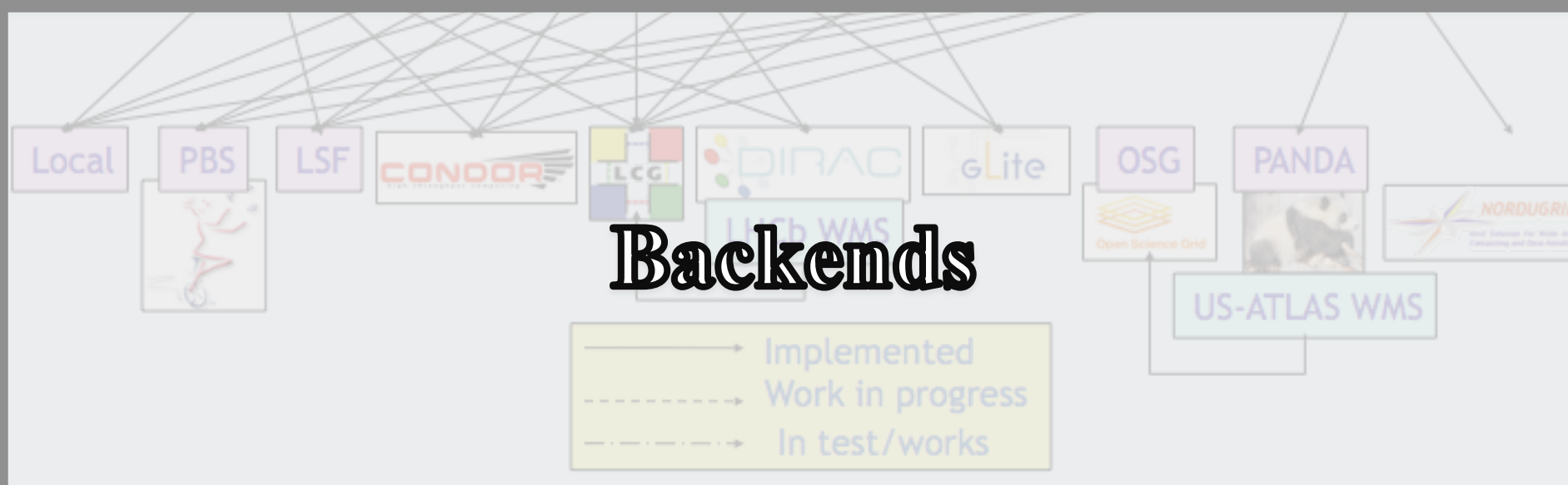
Motivation



Applications



Backends



Motivation



I must learn
many interfaces

How to configure
my applications?



Do I get a consistent view
on all my jobs?



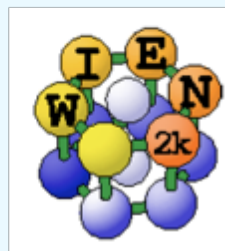
Motivation



Applications



- Originally design to meet the needs of the ATLAS and LHCb for a Grid user interface
- configuring and running applications based on the Gaudi / Athena framework common to the two experiments



Why We Need Ganga



without Ganga

```
Executable = "echo";  
StdOut = "stdout";  
StdErr = "stderr";  
Arguments = "hello world"; write JDL  
... ..  
voms-proxy-init -voms atlas  
glite-wms-job-submit ... hello.jdl  
glite-wms-job-status ... xxxx:yyyy:zzzz  
glite-wms-job-outputs ... xxxx:yyyy:zzzz
```

The 'without Ganga' workflow is a linear sequence of five steps, each in a blue box. The first box contains JDL code and the instruction 'write JDL'. A yellow arrow points down to the second box, which contains the command 'voms-proxy-init -voms atlas' and the instruction 'start grid proxy'. Another yellow arrow points down to the third box, which contains 'glite-wms-job-submit ... hello.jdl' and 'submit job'. A third yellow arrow points down to the fourth box, which contains 'glite-wms-job-status ... xxxx:yyyy:zzzz' and 'check job status'. A final yellow arrow points down to the fifth box, which contains 'glite-wms-job-outputs ... xxxx:yyyy:zzzz' and 'get job outputs'.

with Ganga

```
j = Job(backend=LCG(middleware='glite'))  
j.application = Executable()  
j.application.exec = 'echo'  
j.application.args = ['hello world']  
j.submit()  
j.status
```

- 📖 Ganga knows about the application setup/preparation in terms of backend jobs
- 📖 Ganga does bookkeeping for you
- 📖 Ganga offers programming interface natively in python

What is Ganga



- **Ganga is an easy-to-use frontend for job definition and management**
- **Simplified use of Grid**
- **allows trivial switching between testing on a local batch system and large-scale processing on Grid resources**
- **implemented in [Python](#)**
- **The latest version: GANGA 5.4.0**
 - released 28 Oct, 2009



The Purpose of Ganga



- **Easy for users to create, submit and monitor the progress of jobs**
- **Move transparently from different resources**
- **Keeps track of all jobs and their status through a repository that archives all information between independent Ganga sessions**
- **Simplify the progression from**
 - Rapid prototyping on a local computer
 - Small-scale tests on a local batch system
 - The analysis of a large dataset using Grid resources



Features of Ganga

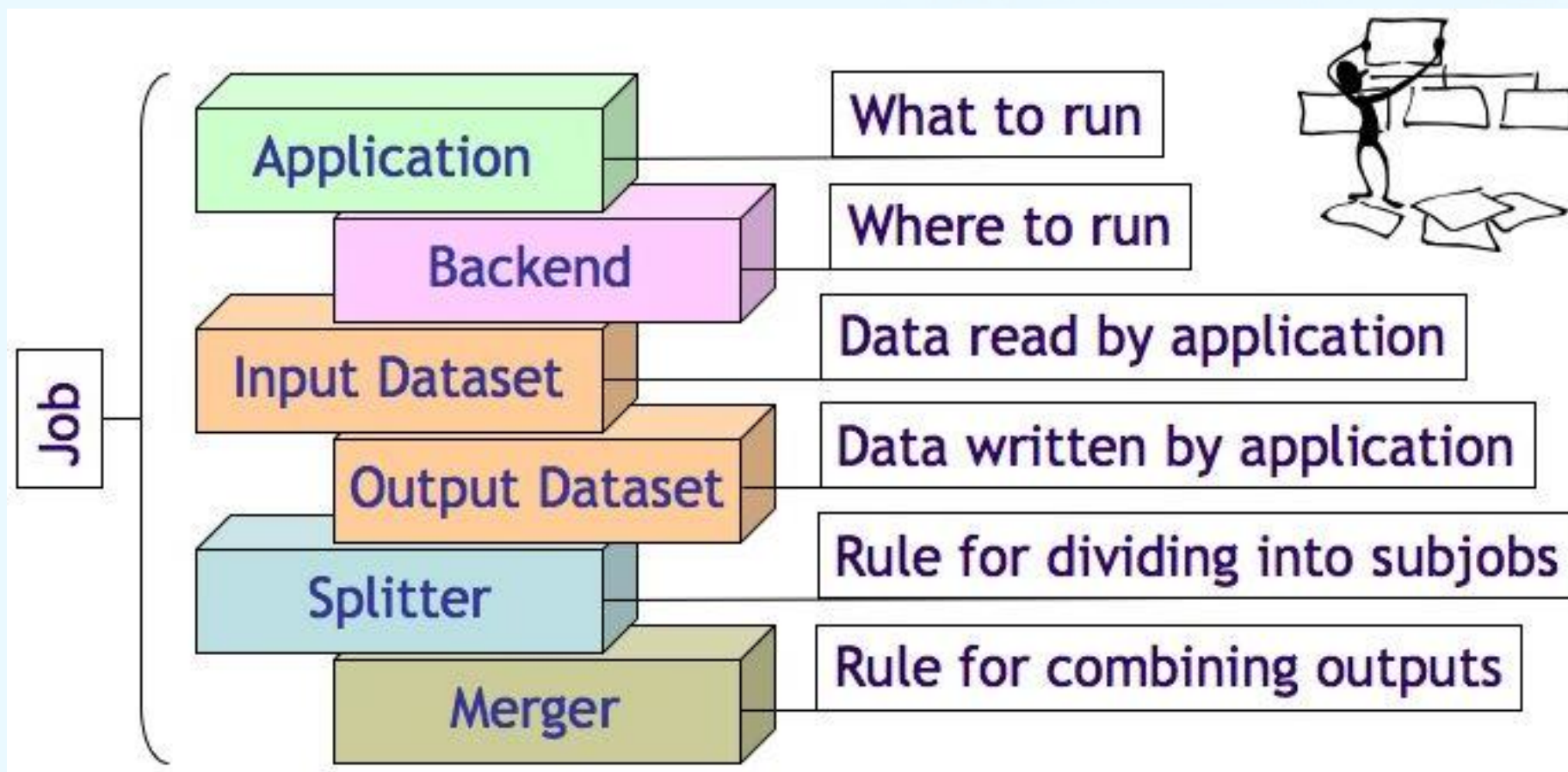
- **A user- and application-oriented layer above existing job submission and management technologies**
- **Interchangable and Interoperability**
 - Ex. Globus, Condor, UNICORE or gLite
- **Encapsulate all low-level setup of the application**
- **Hidden the heterogeneity of backends and data access**



Job Component

- **The code to execute**
- **Input data for processing**
- **Data produced by the application**
- **The specification of the required processing environment**
- **Post-processing tasks**
- **Metadata for bookkeeping**

Ganga Job Components





Ganga Job Components



- **Application**
 - The type of computational task
 - Define the software to be run
- **Backend**
 - Define the processing system to be used
 - Including Localhost, batch systems(PBS), Load Sharing Facility(LSF), Sun Grid Engine, Condor, Grid Systems

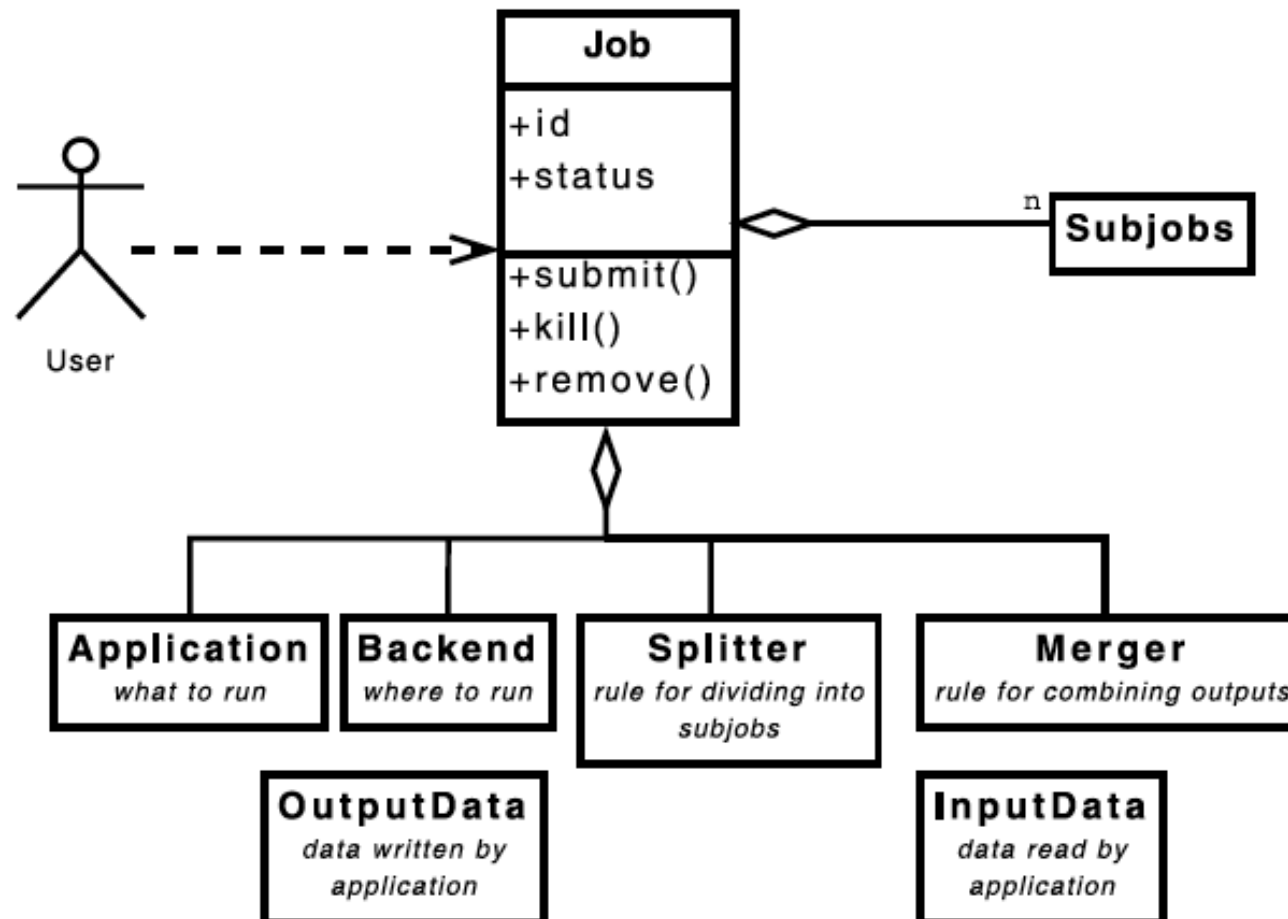


Ganga Job Components



- **Dataset**
 - Uniquely identify a particular collection of data
 - Provide methods for obtaining information about it, such as its location and size.
 - The Descriptions of Data Collections
 - Vary from different problem domains
- **Splitter**
 - Specify the number of subjobs to be created
- **Merger**
 - Allows for the aggregation of subjob outputs

The Component Structure of A Job



To Form a Complete Job



- **Application**

- exe: the path to an executable binary or script
- args: a list of arguments to be passed to the executable
- env: a dictionary of environment variables and the values they should be assigned before the executable is run

```
j = Job()  
j.application = Executable(exe=File('/opt/anotherscript'), args=['-d',File('/  
etc/x')])  
j.submit()
```


To Form a Complete Job



- **Backend**

- Parameters describing the behaviour of a processing system
- Differs among different backend systems
- Methods for job executions, retrieval of jobs status

In [86]:plugins("backends")

Out[86]: ['LSF', 'LCG', 'Dirac', 'gLite', 'PBS', 'Condor', 'Local']

To Form a Complete Job



- **Dataset**
 - Datasets
 - the files or databases stored externally
 - Sandbox
 - consists of files transferred from the user's file system together with the job
 - Sandbox mechanism handles small files (typically up to 10MB)

```
j.inputsandbox = [File(extra_file')]
j.inputsandbox = ['extra_file']
```

```
j.outputsandbox = ['b.dat','a*.txt']
```

To Form a Complete Job



- **Splitter**

- ArgSplitter:

- Deals with executing the same task many times over with changing arguments each time

- args: list of sets of arguments to be passed to an application

```
In [21]:s = ArgSplitter(args=[['A'],['B'],['C']])
```

```
In [22]:j = Job(splitter=s)
```

```
In [23]:j.submit()
```

```
Ganga.GPIDev.Lib.Job           : INFO    submitting job 164
```

```
Ganga.GPIDev.Adapters         : INFO    submitting subjob 16400001
```

```
Ganga.Lib.Localhost           : INFO    job 16400001 submitted
```

```
Ganga.GPIDev.Adapters         : INFO    submitting subjob 16400002
```

```
Ganga.Lib.Localhost           : INFO    job 16400002 submitted
```

```
Ganga.GPIDev.Adapters         : INFO    submitting subjob 16400003
```

```
Ganga.Lib.Localhost           : INFO    job 16400003 submitted
```

```
Out[23]: 1
```

To Form a Complete Job



- **Merger**

- Combing data in a particular format
- (ex. Text strings or data representing histograms)
- TextMerger:
 - concatenate the files of standard output and error returned by a set of subjobs
- RootMerger:
 - sums the histograms produced in ROOT format

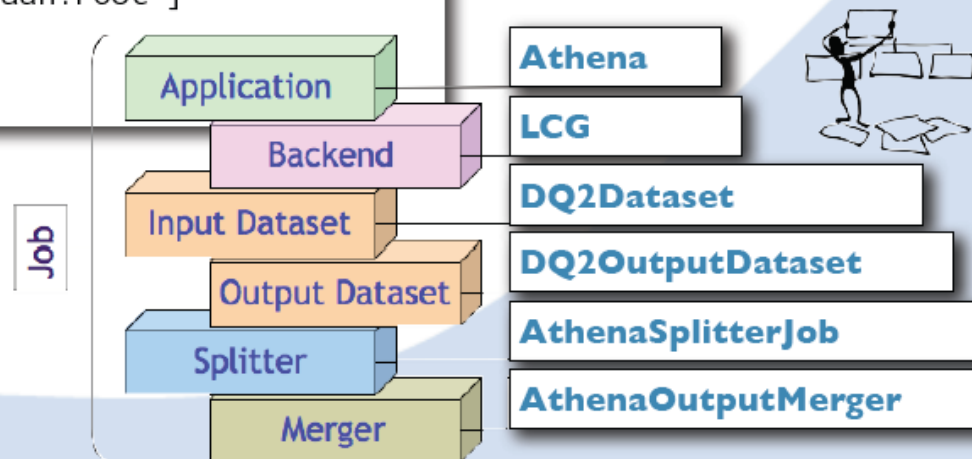
```
In [4]:j.merger=TextMerger()  
In [5]:j.merger.files=['stdout']  
In [6]:j.merger.ignorefailed = True  
In [7]:j.submit()
```

```
In [17]:j.peek()  
total 3.0K  
-rw-r--r--  1 moscicki sf  903 Jun 11  
13:08 stdout.merge_summary  
-rw-r--r--  1 moscicki sf 1.2K Jun 11  
13:08 stdout
```

Job Representation - Example



```
j = Job()
j.application=Athena()
j.application.prepare(athena_compile=False)
j.application.option_file='$HOME/athena/12.0.5/InstallArea/jobOptions'
j.splitter=AthenaSplitterJob()
j.splitter.numsubjobs = 3
j.merger=AthenaOutputMerger()
j.inputdata=DQ2Dataset()
j.inputdata.dataset='csc11.005145.PythiaZmumu.recon.AOD.v11004103'
j.inputdata.match_ce=True
j.outputdata=DQ2OutputDataset()
j.outputdata.outputdata=['AnalysisSkeleton.aan.root']
j.backend=LCG()
j.submit()
```



Job Representation - Example



```
j = Job()
j.application=Athena()
j.application.prepare(athena_compile=False)
j.application.option_file=['$HOME/Atlas/testarea/14.2.10/PhysicsAnalysis/
AnalysisCommon/UserAnalysis/run/AnalysisSkeleton_topOptions.py']
j.application.max_events='100'
```

Application

```
j.inputdata=DQ2Dataset()
j.inputdata.dataset='trig1_misal1_rh12005322.PythiaVBFH170wwll.recon.AOD.
v13003003 tid017852'
```

Dataset

```
j.outputdata=DQ2OutputDataset()
j.outputdata.outputdata=['AnalysisSkeleton.aan.root' ]
```

Dataset

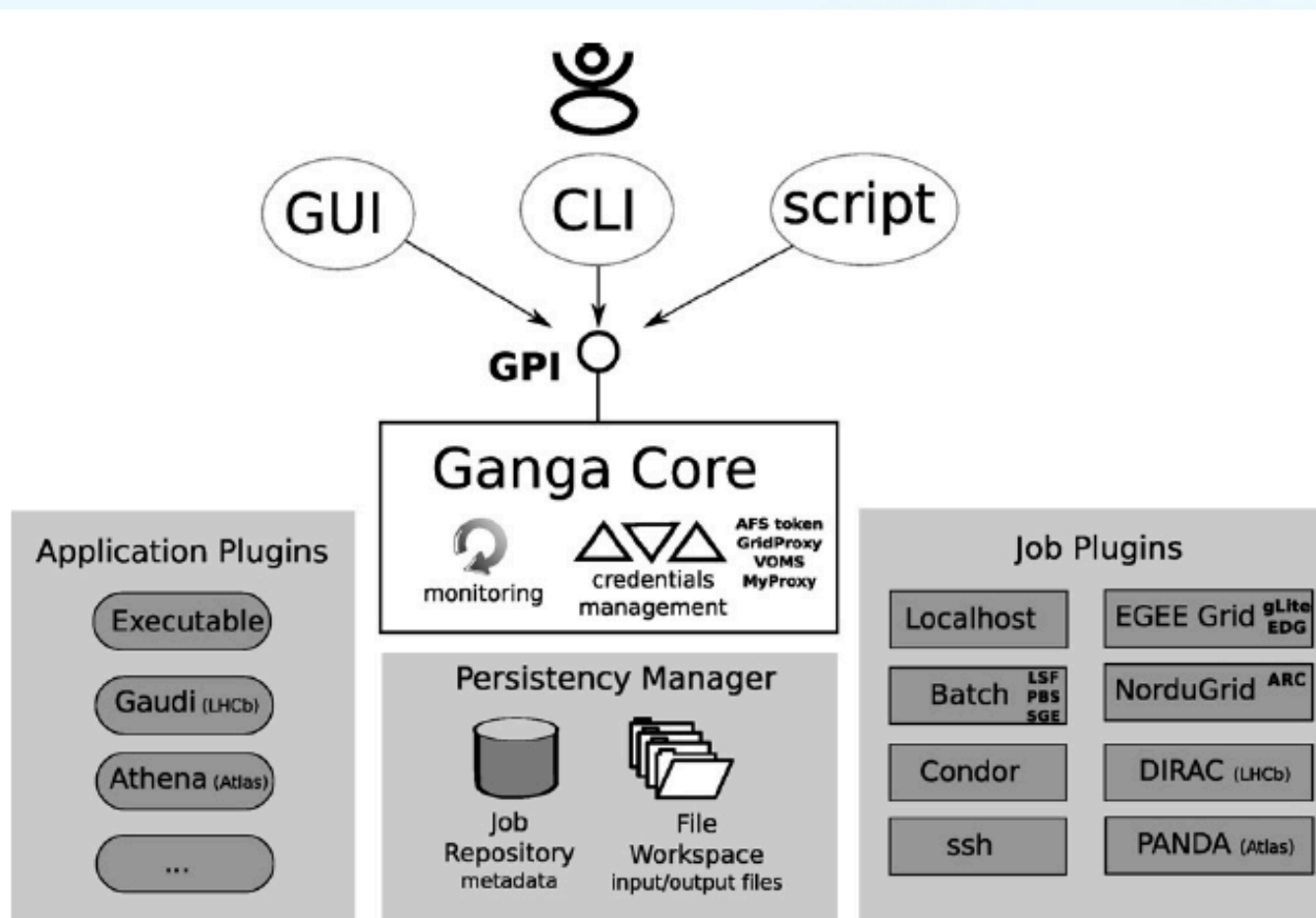
```
j.splitter=DQ2JobSplitter()
j.splitter.numsubjobs=3
j.merger=AthenaOutputMerger()
```

Splitter & Merger

```
j.backend=LCG()
j.backend.requirements.cloud='DE'
j.submit()
```

Backend

Ganga Architecture



Ganga Interfaces



- **Built on top of the Ganga Public Interface (GPI)**
- **A Text-based Command Line in Python reference**
- **A File-based Scripting Interface**
- **Graphical User Interface (GUI)**

A File-based Scripting Interface



- From the command line, a script `myScript.py` can be executed in the Ganga environment using:
 - `Shell> ganga myScript.py`
- `myScript.py`

```
import os
dir=os.environ['HOME']+'/mydir'
file=os.listdir(dir)
for line in file:
    print line
    j = Job(Executable(exe=File(dir+'/'+line)))
    j.submit
```

An example of a ganga script

Ganga GUI



Scriptor

Job details

Logical Folders

Job Monitoring

Job builder

Log window

There is also a scripting interface (like pAthena)
I will use the line mode.....

Ganga GUI



Job Folders View Help

Scriptor Log Job Builder

Jobs

id	status	name	application	exe filename	backend	bc
0	failed	G4Bnchy	Executable	File	LCG	N3
1	completed		Executable	echo	LCG	ht
2	completed	DIANEWorkerAgent	Executable	File	LCG	ht
3	completed	DIANEWorkerAgent	Executable	File	LCG	ht
4	completed	DIANEWorkerAgent	Executable	File	LCG	ht
11	completed	DIANEWorkerAgent	Executable	File	LCG	ht
12	completed	DIANEWorkerAgent	Executable	File	Local	25
13	completed		Executable	echo	Local	12
14	new		Executable	echo	Local	-1
15	new		Executable	echo	Local	-1
16	failed		Executable	echo	Local	93
17	failed		Executable	echo	Local	15
18	completed		Executable	echo	Local	19

Job Details

```
Job (
  status = 'completed',
  name = '',
  inputdir = '/afs/cern.ch/user/m/moscicki/ganga',
  outputdir = '/afs/cern.ch/user/m/moscicki/ganga',
  outputsandbox = [],
  id = 1,
  info = JobInfo (
    submit_counter = 1
  ),
  inputdata = None,
  merger = None,
  inputsandbox = [],
  application = Executable (
    exe = 'echo',
    env = {}
  )
)
```

removing job 5
removing job 6
removing job 7
removing job 8
removing job 9
removing job 10



Ganga Public Interface (GPI)



- **User-level Interface separated from low-level, internal API**
- **Templates of Job Configurations**
 - Frequently used job configurations
- **Hierarchical JobTree**
 - Jobs can be labelled and organised in a hierarchical jobtree

- **Credentials Management**
 - User credentials, including classic Grid proxies with extensions for VOMS
 - Renew and destroy the credentials using GPI
 - Multiple security models
- **Monitoring**
 - Internal Monitoring
 - External Monitoring

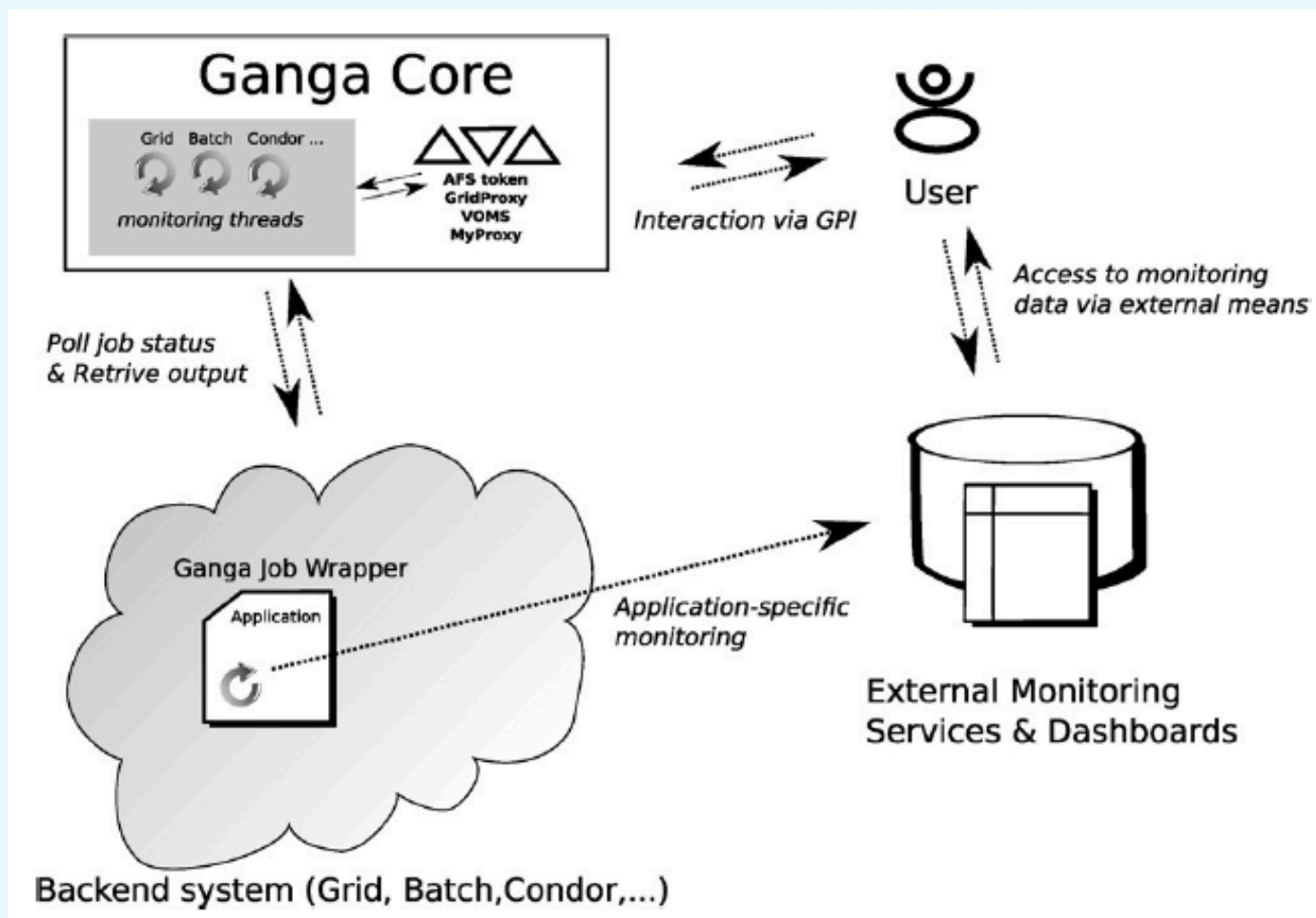


Ganga Core – Job Monitoring



- **Internal Monitoring**
 - Polling job status for varying backends
 - The remaining validity of authentication credentials
- **External Monitoring**
 - Dynamically adding third-party monitoring sensors
 - Monitoring sensor
 - Client side
 - Remote environment
 - Allow collection of both
 - generic execution information
 - Application-specific data

Core - Job Monitoring





Persistence Manager



- **Job Repository Database**
 - Job Persistence and Objects
 - Job Bookkeeping and Metadata
 - Local and Remote Repository
 - Schema Migration
- **File Workspace**
 - The Input and Output files associated with the Jobs

Application Plugins



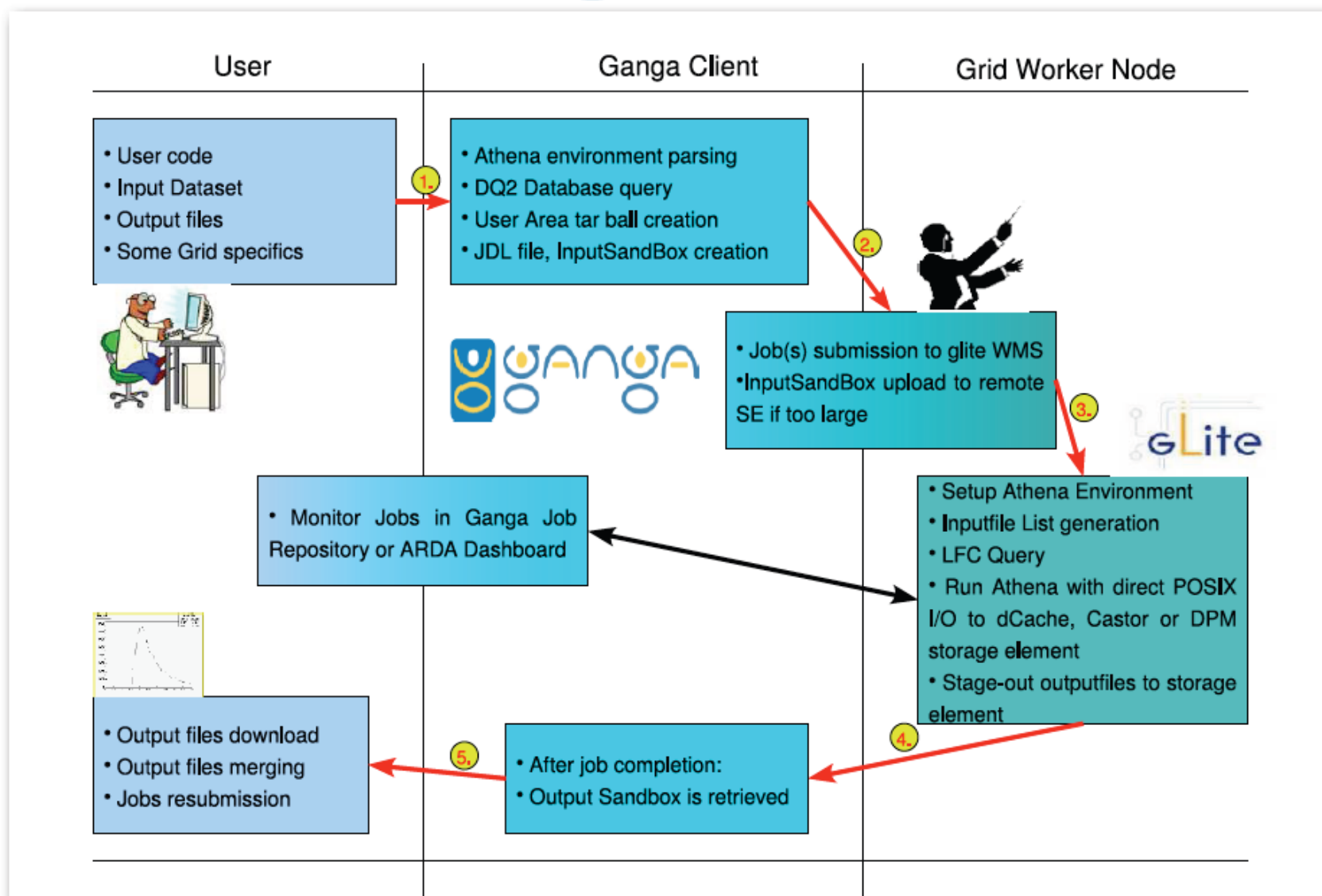
- **Provides a configuration object which contains backend- independent information:**
 - The application to be run
 - The user code to be executed
 - The values to be assigned to any configurable parameters
 - The data to be processed

Job Plugins



- **Accepts an input configuration object and calls the application runtime handler responsible for the backend-specific part of the application configuration and management:**
 - Job splitting
 - Packaging of user code
 - Submitted jobs to the backend
 - Monitoring job progress
 - Retrieving output files when jobs complete

Ganga Workflow



Python script – Running Athena



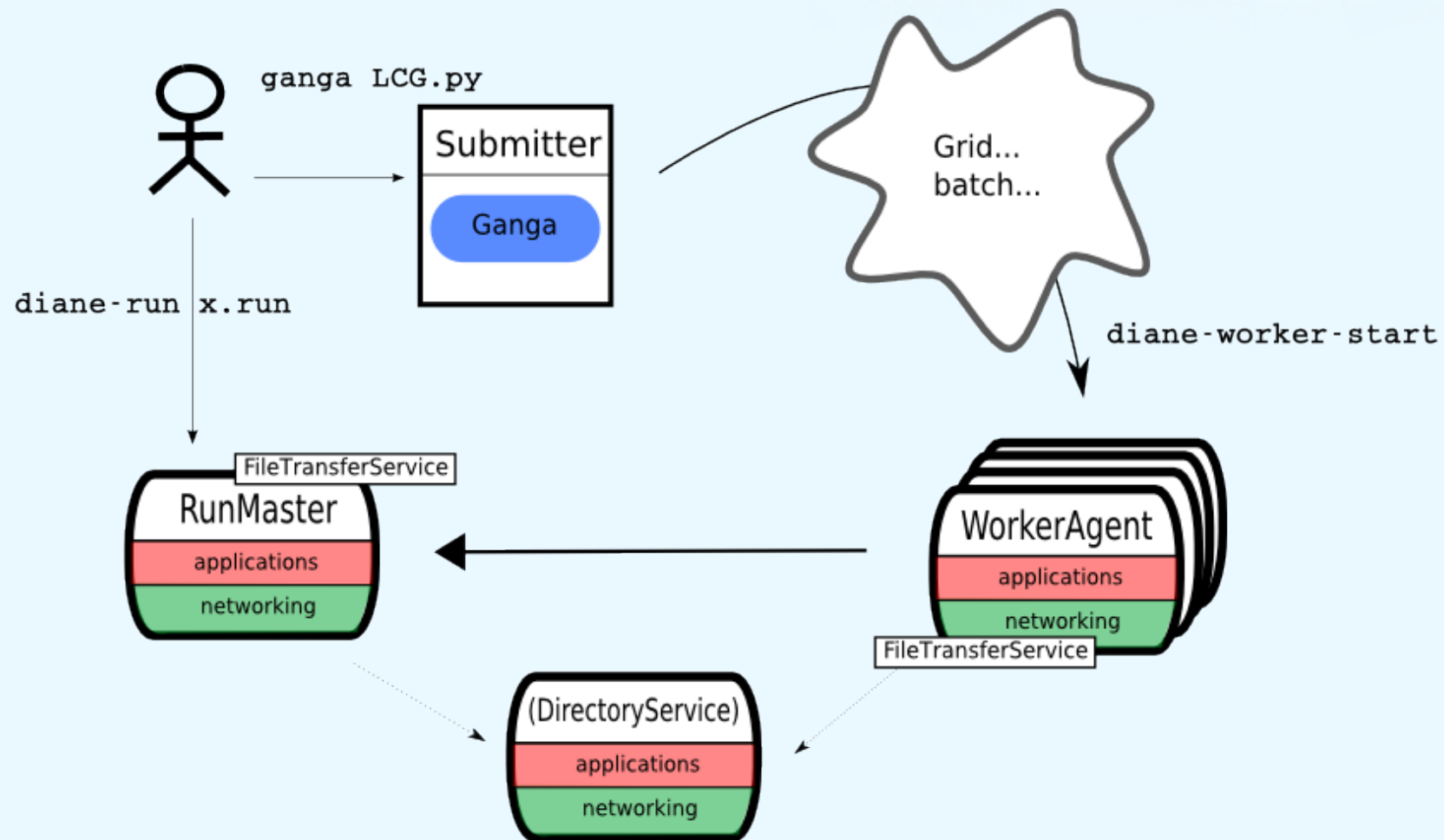
```
j = Job()
j.application=Athena()
j.application.exclude_from_user_area=["*.o","*.root","*.exe"]
j.application.prepare(athena_compile=True)
j.application.atlas_release="14.2.10"
j.application.option_file=['my_jobOption'] <= jobOption filename (absolute path) //
j.application.max_events='100'
j.inputdata=DQ2Dataset()
j.inputdata.dataset="mc08.007081.singlepart_gamma_E10.recon.AOD.e339_s439_r462_tid023328"
j.inputdata.number_of_files=1
<= if you want to analyze a specified number of files
j.inputdata.type='DQ2_LOCAL'
file directly on SE // 'DQ2_DOWNLOAD' to force copy of file to WN )
j.outputdata=DQ2OutputDataset()
j.outputdata.outputdata=['recon.CBNT.pool.root']
j.outputdata.location="IN2P3-LAPP_LOCALGROUPDISK"
j.splitter=DQ2JobSplitter() <= if you want to analyze complete dataset
j.splitter.numsubjobs=6 <= (specify the number of subjobs)
j.merger=AthenaOutputMerger()
j.backend=LCG() j.backend.requirements.cloud='FR' //
j.backend.requirements.sites= ['LYON_MCDISK'] (dq2-ls -r command) j.submit()
```


Ganga & DIANE

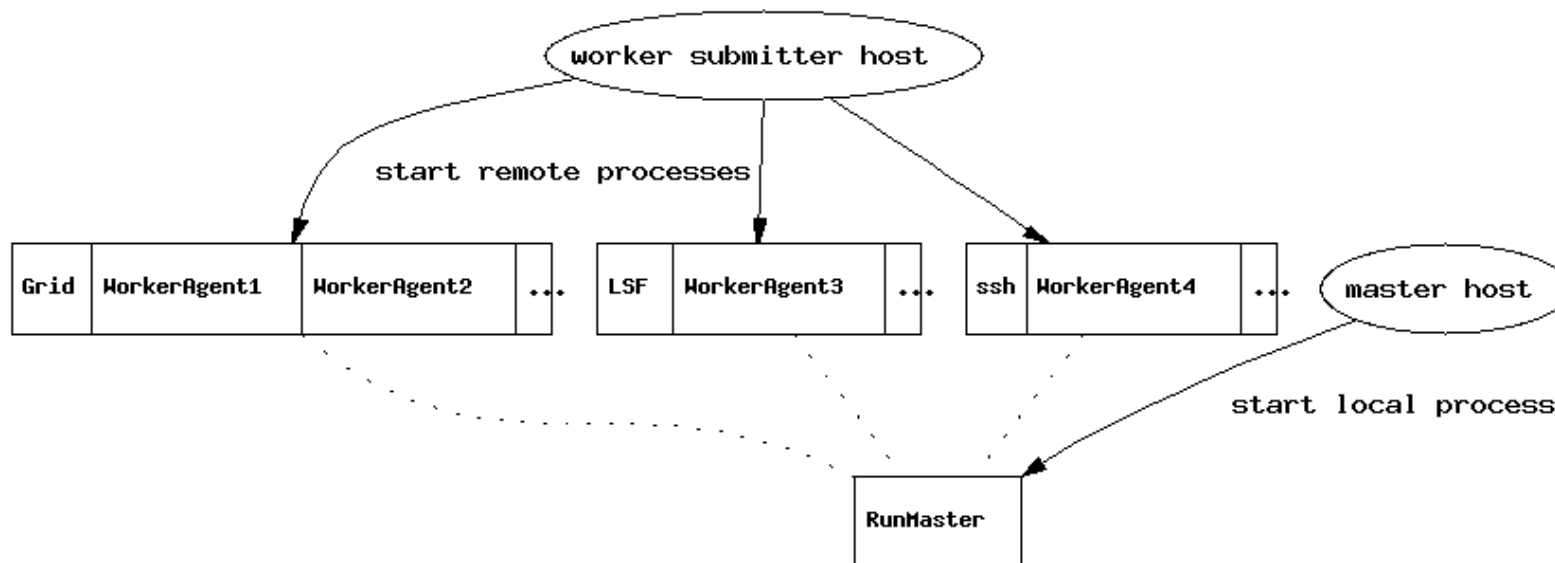


- **DIANE**
 - A lightweight agent-based scheduling layer on top of the Grid
 - Master & Worker Model
- **Ganga used in combination with DIANE**
 - DIANE works as an overlay scheduling system
 - Ganga works as a well-structured job management tool
- **DIANE worker agents are executed as Ganga jobs**

Ganga Submitter in DIANE Architecture



Ganga Job Submission tool in DIANE



Use Case

Ganga Interfaced to DIANE



Framework

- **Drug Discovery, Auto-docking Tool developed by ASGC, TW**
 - Ganga embedded in web-based services
 - This tool is for analysis of candidate drugs against avian flu
- **Ganga as a job management component embedded in DIANE, with the application**

Drug Discovery Tool with Ganga



Virtual Screening Service

Target Ligand Docking Parameter

Default Targets

Target : DC2_T01IAN

Name	Owner
DC2_T01IAN	DEFAULT
DC2_T02IAN	DEFAULT
DC2_T03IAN	DEFAULT
DC2_T04IAN	DEFAULT
DC2_T05IAN	DEFAULT
DC2_T06IAN	DEFAULT
DC2_T07IAN	DEFAULT
DC2_T08IAN	DEFAULT
Dengue_NS3	DEFAULT

Visualize

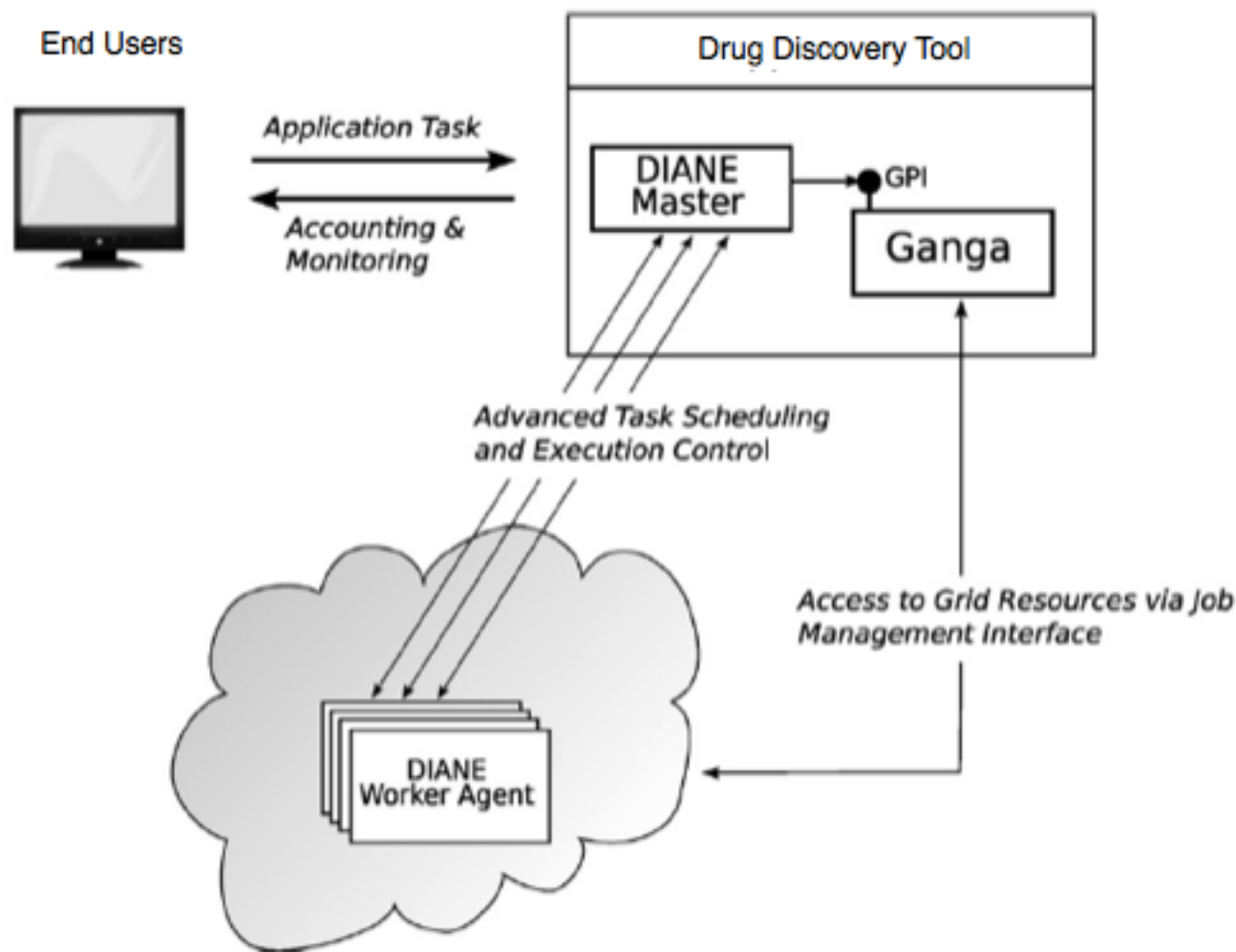
Target Structure

[TRP]189:B.2HB #1618

DC2_T01IAN

Job Description : Resource Domain : GRID Worker Number : 10 Submit

Drug Discovery Tool with Ganga



Ganga as a job management component embedded in DIANE, with the application,
Drug Discovery Tool

References



- **Ganga Home**
 - <http://ganga.web.cern.ch/ganga/index.php>
- **Ganga User Guide**
 - <http://ganga.web.cern.ch/ganga/user/index.php>
- **Ganga GPI Reference**
 - <http://ganga.web.cern.ch/ganga/release/4.3.2/reports/html/Manuals/GangaTutorialManual.html>

~ The End ~

