

# Advanced use of Workload Management System

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Tutorial CYCLOPS





- **MPI Jobs**
- **WMPProxy Overview**
  - **Special Jobs**
    - DAG jobs
    - Job collections
    - Parametric jobs





- Execution of parallel jobs is an essential issue for modern informatics and applications.
- Most used library for parallel jobs support is **MPI (Message Passing Interface)**
- At the state of the art, parallel jobs can run **inside single Computing Elements (CE) only**;
  - several projects are involved into studies concerning the possibility of executing parallel jobs on Worker Nodes (WNs) belonging to different CEs.
- The source code must have been compiled with **mpicc** libraries





# MPI JDL

- ```
> $ cat mpi.jdl  
[  
  Type = "Job";  
  #Mandatory  
  JobType = "MPICH";  
  #Bash script containing invocation to MPI  
  #interpreter, mpirun  
  Executable = "MPIscript.sh";  
  #The number of CPU that will be used  
  NodeNumber = 2;  
  StdOutput = "cpi.out";  
  StdError = "cpi.err";  
  InputSandbox = {"cpi","MPIscript.sh"};  
  OutputSandbox = {"cpi.err","cpi.out"};  
  RetryCount = 3;  
]
```





# MPI jobs @ work

- Login to the UI :
  - `ssh bolognaXX@glite-tutor.ct.infn.it`
- initialize a proxy if you haven't it
  - `voms-proxy-init --voms gilda`
- `cd examples`
- `edg-job-submit mpi.jdl`
- Monitor job status and finally get the output using the job id given by the submit command





# Workload Manager Proxy





- **WMPProxy (Workload Manager Proxy)**

- is a new service providing access to the gLite Workload Management System (WMS) functionality through a simple Web Services based interface.
- has been designed to handle a large number of requests for job submission
  - gLite 3.0=> ~180 secs for 500 jobs
  - goal is to get in the short term to ~60 secs for 1000 jobs
- it provides additional features such as *bulk submission* and the support for *shared and compressed* sandboxes for *compound jobs*.
- It's the natural replacement of the NS in the passage to the *SOA approach*.





- **Support for new types strongly relies on newly developed JDL converters and on the DAG submission support**
  - all JDL conversions are performed on the server
  - a single submission for several jobs
- **All new request types can be monitored and controlled through a single request id**
  - each sub-jobs can be however followed-up and controlled independently through its own id
- **“Smarter” WMS client commands/API**
  - allow submission of DAGs, collections and parametric jobs exploiting the concept of “shared sandbox”
  - allow automatic generation and submission of collections and DAGs from sets of JDL files located in user specified directories on the UI





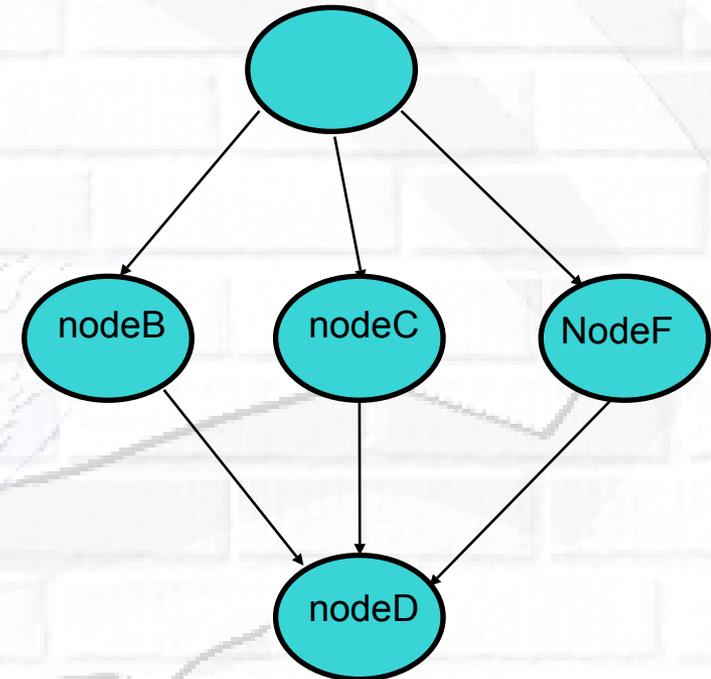
## Special Jobs

- **DAG**
- **Job Collection**
- **Parametric jobs**





- A DAG job is a set of jobs where input, output, or execution of one or more jobs can depend on other jobs
- Dependencies are represented through **Directed Acyclic Graphs**, where the nodes are jobs, and the edges identify the dependencies





- Type = "DAG" → *Mandatory*
- VirtualOrganisation = "yourVO" → *Mandatory*
- Max\_Nodes\_Running = int >0 → *Optional*
- MyProxyServer = "... " → *Optional*
- Requirements = "... " → *Optional*
- Rank = "... " → *Optional*
- InputSandbox =  more later! → *Optional*
- ~~• OutSandbox = "... " → *Optional*~~
- Nodes = nodeX  more later! → *Mandatory*
- Dependencies  more later! → *Mandatory*



The *Nodes* attribute is the core of the DAG description;

```
....  
Nodes = [ nodefilename1 = [...]  
           nodefilename2 = [...]  
           .....  
           dependencies = ...  
         ]
```

```
Nodefilename1 = [ file = "foo.jdl"; ]  
Nodefilename2 =  
  [ file = "/home/vardizzo/test.jdl";  
    retry = 2;      ]
```

```
Nodefilename1 = [  
  description = [ JobType = "Normal";  
                  Executable = "abc.exe";  
                  Arguments = "1 2 3";  
                  OutputSandbox = [...];  
                  InputSandbox = [...];  
                  ..... ]  
  retry = 2;  
  ]
```

# Attribute: Dependencies



- It is a list of lists representing the dependencies between the nodes of the DAG.

```
....  
Nodes = [ nodefilename1 = [...]  
          nodefilename2 = [...]  
          .....  
          dependencies = ...  
        ]
```



```
dependencies =  
    {nodefilename1, nodefilename2}
```



**MANDATORY : YES!**

```
dependencies = {};
```

```
{ nodefilename1, nodefilename2 }
```

```
{ { nodefilename1, nodefilename2 }, nodefilename3 }
```

```
{ { { nodefilename1, nodefilename2 }, nodefilename3 }, nodefilename4 }
```



```
[
  type = "dag";
  max_nodes_running = 4;
  nodes = [
    nodeA = [
      file = "nodes/nodeA.jdl" ;
    ];
    nodeB = [
      file = "nodes/nodeB.jdl" ;
    ];
    nodeC = [
      file = "nodes/nodeC.jdl" ;
    ];
    nodeD = [
      file = "nodes/nodeD.jdl";
    ];
    dependencies = {
      {nodeA, nodeB},
      {nodeA, nodeC},
      { {nodeB,nodeC}, nodeD }
    }
  ];
]
```

Node description could also be done here, instead of using separate files



- A job collection is a set of independent jobs that user wants to submit and monitor via a single request
- Jobs of a collection are submitted as DAG nodes without dependencies
- JDL is a list of classad, which describes the subjobs

```
[  
    Type = "collection";  
    VirtualOrganisation = "gilda";  
    nodes = {  
        [ <job descr 1 >],  
        [ <job descr 2 >],  
        ...  
    };  
]
```



# 'Scattered' Input Sandboxes

- **Input Sandbox can contain**
  - file paths on the UI machine (i.e. the usual way)
  - pointer to other files within the DAG/collection
  - URI pointing to files on a remote gridFTP/HTTPS server

```
InputSandbox = {  
    "gsiftp://neo.datamat.it:2811/var/prg/sim.exe",  
    root.nodes.nodeA.description.OutputSandbox[0],  
    "file:///home/pacio/myconf" };
```

- **Only local files (`file://`) are uploaded to the WMS node**
- **File pointed by URIs are directly downloaded on the WN by the JobWrapper just before the job is started**





# 'Scattered' Output Sandboxes

- JDL is enriched with attributes, specifying the destinations for the files listed within OutputSandbox attribute list

```
OutputSandbox = {      "jobOutput","run1/event1",  
                      "jobError"                };  
  
OutputSandboxDestURI = {  
    "gsiftp://matrix.datamat.it/var/jobOutput",  
    "https://grid003.ct.infn.it:8443/home/cms/event1",  
    "gsiftp://matrix.datamat.it/var/jobError"    };
```

- A base URI to be applied to all sandbox files can also be specified

```
OutputSandboxBaseDestURI = "gsiftp://neo.datamat.it/home/run1/";
```

- Files are copied when the job has completed execution by the JobWrapper to the specified destination without transiting on the WMS node

# Job collection example



```
[
  type = "collection";
  InputSandbox = {"date.sh"};
  RetryCount = 3;
  nodes = {
    [
      file = "jobs/job1.jdl" ;
    ],
    [
      [
        Executable = "/bin/sh";
        Arguments = "date.sh";
        StdOutput = "date.out";
        StdError = "date.err";
        OutputSandbox = {"date.out", "date.err"};
      ]
    ],
    [
      file = "jobs/job3.jdl" ;
    ]
  };
]
```

All nodes will share  
this Input Sandbox





- A parametric job is a job where one or more of its attributes are parameterized
- Values of attributes vary according to a parameter

```
[  
  JobType = "Parametric";  
  Executable = "/bin/sh";  
  Arguments = "md5.sh input_PARAM.txt";  
  InputSandbox = {"md5.sh", "input_PARAM.txt"};  
  StdOutput = "out_PARAM.txt";  
  StdError = "err_PARAM.txt";  
  Parameters = 4;  
  ParameterStart = 1;  
  ParameterStep = 1;  
  OutputSandbox = {"out_PARAM.txt", "err_PARAM.txt"};  
]
```

- Job monitoring / managing is always done through a unique jobID, as if the job was single (see submission of collection)





- Parameter can be also a list of string
- InputSandbox (if present) has to be coherent with parameters

```
[ui-test] /home/giorgio/param > cat param2.jdl  
[  
    JobType = "Parametric";  
    Executable = "/bin/cat";  
    Arguments = "input_PARAM_.txt";  
    InputSandbox = "input_PARAM_.txt";  
    StdOutput = "myoutput_PARAM_.txt";  
    StdError = "myerror_PARAM_.txt";  
    Parameters = {earth,moon,mars};  
    OutputSandbox = {"myoutput_PARAM_.txt"};  
]
```

```
[ui-test] /home/giorgio/param > ls  
inputEARTH.txt  inputMARS.txt  inputMOON.txt  param2.jdl
```



- **gLite 3.0 User Guide**
  - <https://edms.cern.ch/file/722398/1.1/gLite-3-UserGuide.pdf>
- **JDL attributes specification for WM proxy**
  - <https://edms.cern.ch/document/590869/1>
- **WMPProxy quickstart**
  - [http://egee-jra1-wm.mi.infn.it/egee-jra1-wm/wmproxy\\_client\\_quickstart.shtml](http://egee-jra1-wm.mi.infn.it/egee-jra1-wm/wmproxy_client_quickstart.shtml)
- **WMS user guides**
  - <https://edms.cern.ch/document/572489/1>
- **MPI**
  - <http://www-unix.mcs.anl.gov/mpi/>

