

# Reference manuals

# Infinity

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<https://infinity.ncbr.muni.cz>

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# Batch processing

**Batch processing** is the execution of a series of programs (so-called batches) on a computer without the participation of the user. Batches are prepared in advance so that they can be processed without the participation of the user. All input data is prepared in advance in files (scripts) or entered using parameters on the command line. Batch processing is the opposite of interactive processing, where the user provides the required inputs only when the program is running.

## Advantages of batch processing

- sharing computer resources between many users and programs
- postponing batch processing until the computer is less busy
- eliminate delays caused by waiting for user input
- maximizing computer utilization improves investment utilization (especially for more expensive computers)

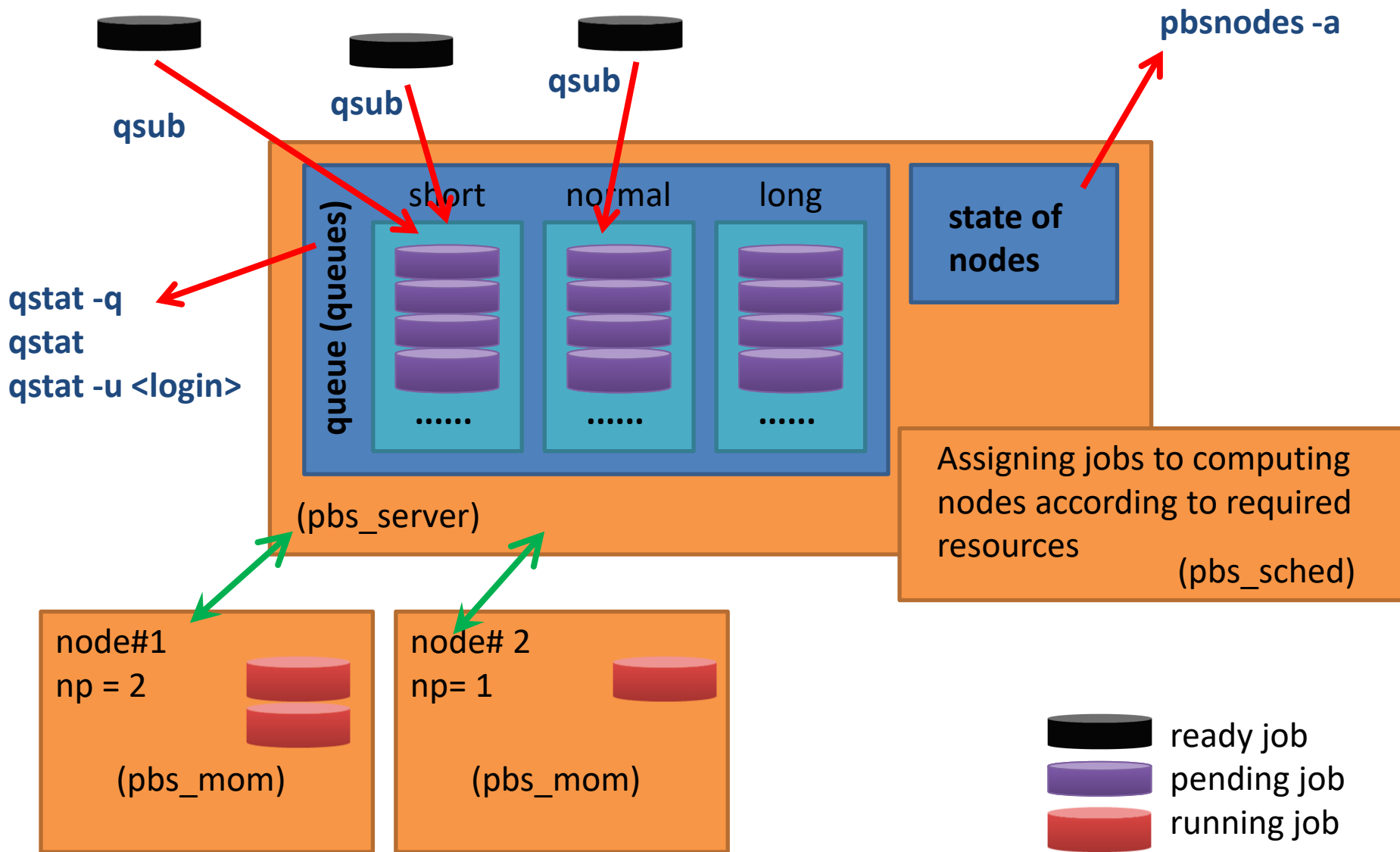
source: [www.wikipedia.cz](http://www.wikipedia.cz), modified

## ➤ PBSPro

<https://www.openpbs.org/>, <https://www.altair.com/>

PBSPro is used as a batch system on our local clusters (WOLF, ...), in MetaCentrum VO, and IT4I.

# Architecture - PBSPro



# PBSPro - commands, job states

<b>qsub</b>	submits job to the batch system
<b>qstat</b>	prints information about the batch system (job list, queue list)
<b>pbsnodes</b>	prints information about computing nodes
<b>qrls</b>	releases job from the state <b>holded</b> (if circumstances allow)

## Job states:

<b>Q</b> (queued)	job is waiting in queue to run on computing node
<b>R</b> (running)	job is running on computing nodes
<b>C</b> (completed)	job has been completed (information about completed tasks is displayed only for a limited time - most often 24 hours)
<b>H</b> (holded)	job has been paused, job can be released with command <b>qrls</b>
<b>E</b> (exiting)	job is ending
<b>F</b> (finished)	job is completed: successful or unsuccessful termination

# Infinity - overview of commands

**Infinity** is a software environment **simplifying management of computational jobs**. It extends functionality of the PBSPro batch environment.

## Software management:

- site            activation of logical computing resources
- software       activation/deactivation of software

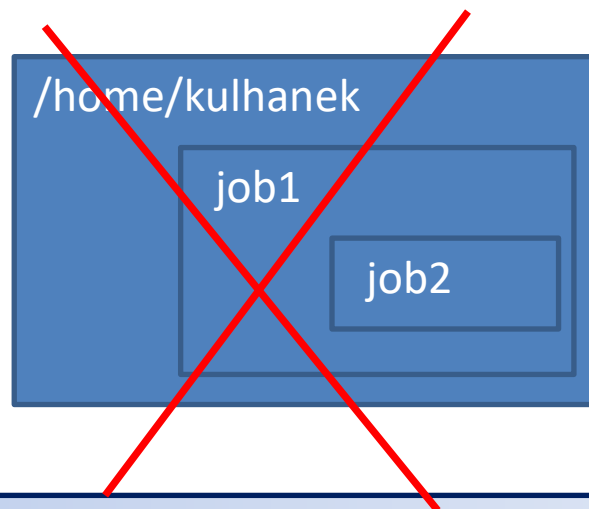
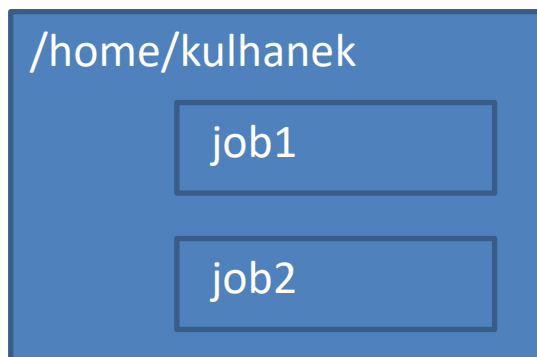
## Task management:

- pqueues       overview of batch system queues available to the user
- pnodes        overview of computing nodes available to the user
- pqstat        overview of all tasks submitted into the batch system
- pjobs         overview of user tasks submitted into the batch system
- psubmit      submitting a job into the batch system
- pinfo         job information
- pgo            logs the user on to the computing node where the task is performed
- psync         manual data synchronization

# Job

## Job **must fulfill** the following conditions:

- each job runs in a separate directory
- all job input data must be in the job directory
- job directories must not be nested
- the progress of the job is controlled by a script or input file (for automatically detected jobs)
- the job script must be written in bash
- absolute paths must not be used in the job script, all paths must be referenced relative to the job directory



# Job script

The job script can be introduced by standard interpreter (**bash**) or special interpreter **infinity-env** which protects the job execution outside the computing node.

The second approach prevents possible damage/overwriting/deletion of already calculated data by accidental re-running of the script.

```
#!/bin/bash
```

```
# script itself
```

```
#!/usr/bin/env infinity-env
```

```
# script itself
```

This is automatically set for auto-detected jobs.

# Submitting a job

The job is submitted **in the job directory** by the **psubmit** command.

```
psubmit destination job [resources]
```

**destination** (where to submit the job) is:

- queue\_name

**job** is either:

- job script name
- input file name for automatically detected jobs (gaussian, orca, etc.)

**resources** are required computational resources. If not specified, 1 CPU is requested.



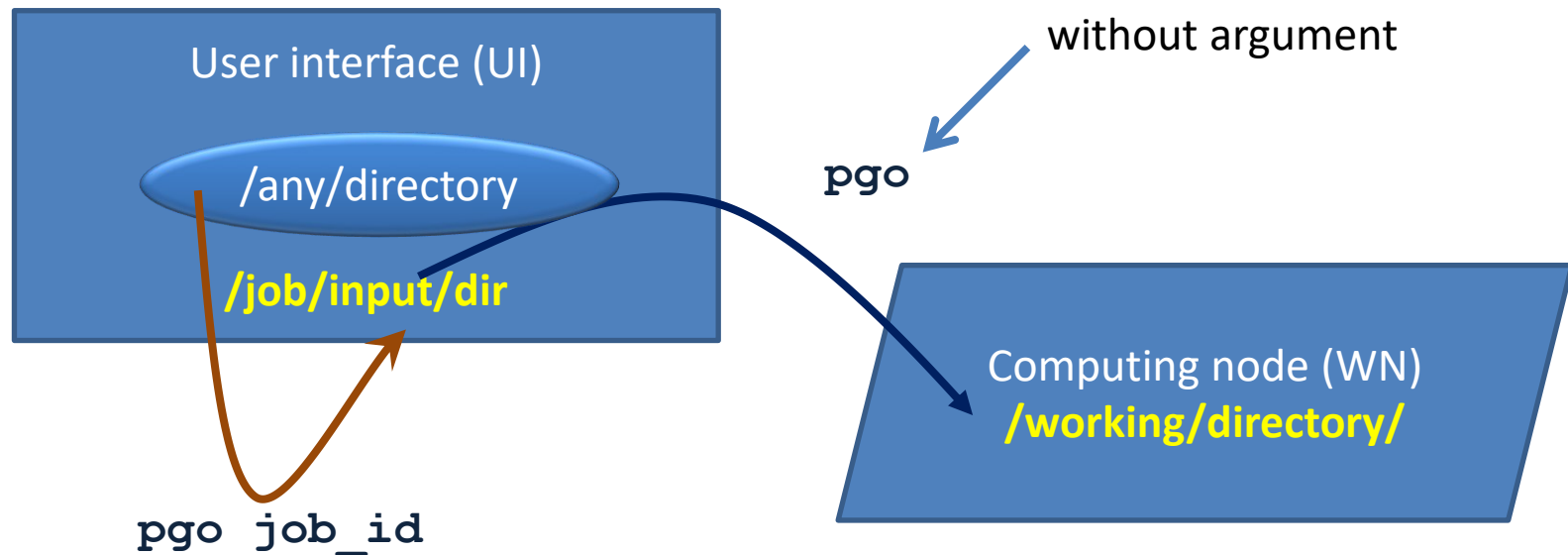
# Resource specifications (selected)

Source	Description
<b>ncpus</b>	total number of CPUs requested
<b>ngpus</b>	total number of GPUs requested
<b>nnodes</b>	number of computational nodes (WN)
<b>mem</b>	total amount of requested memory (CPU), unit mb, gb
<b>walltime</b>	maximum job run time
<b>workdir</b>	type of working directory on WN
<b>place</b>	method of occupying computing nodes
<b>props</b>	required properties of computational nodes

# Monitoring progress of job

You can use command **pinfo** to monitor the progress of the job which is run either in the job input directory or in the working directory on the computing node. Other monitoring possibilities are offered by the **pjobs** and **pqstat** commands.

If the job is running on a computing node, you can use the **pgo** command, which logs the user on to the computing node and changes the current directory to the job working directory.



Monitoring the job in the terminal.

# Service files

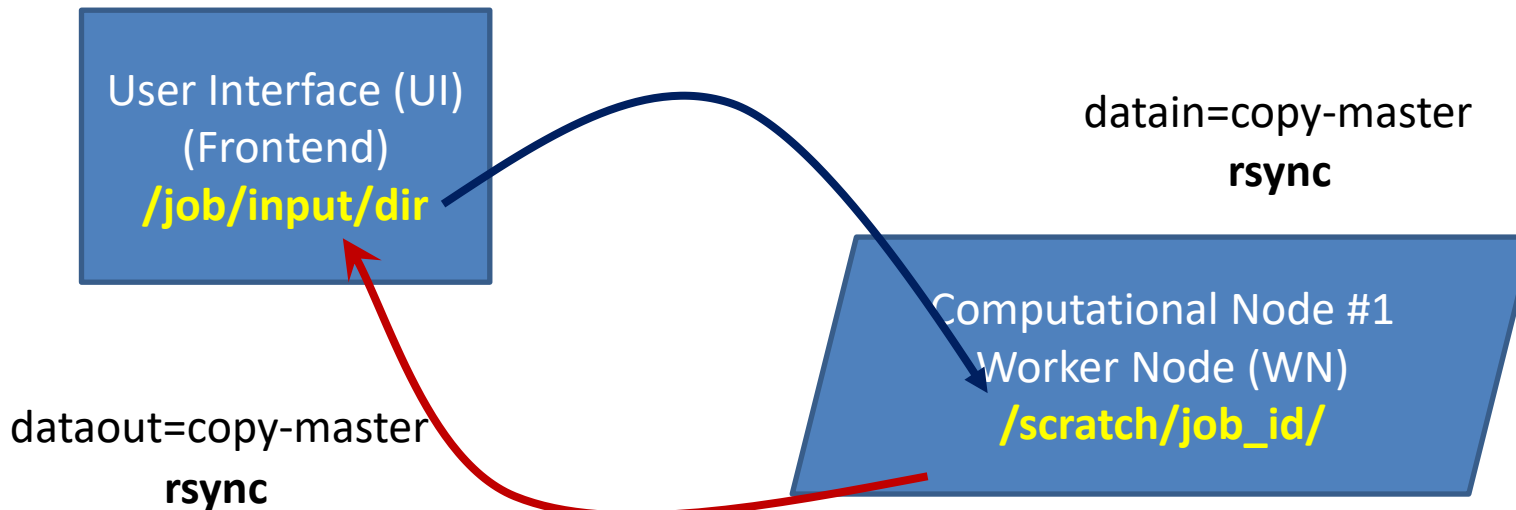
In the job directory, service files are created when the job is submitted into the batch system, during the job life, and after its completion. Their meaning is as follows:

- \*.info control file with information about the progress of the job
- \*.infex custom script (wrapper), which is run by the batch system
- \*.infout standard runtime output of \*.infex script, **must be analyzed when the task terminates abnormally**
- \*.nodes list of nodes reserved for the job
- \*.mpinodes list of nodes reserved for the job in format for OpenMP
- \*.gpus list of GPU cards reserved for the job
- \*.key unique job identifier
- \*.stdout **standard output from a job script**

# Data synchronization

## Default operating mode

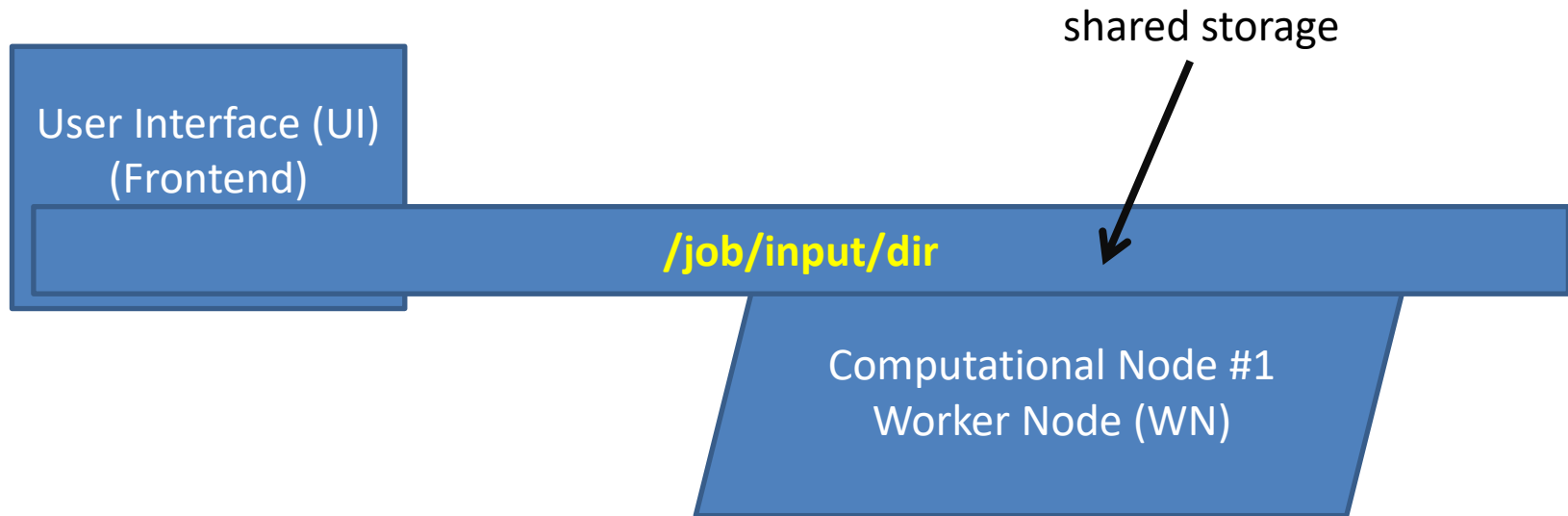
Source	Meaning
<code>workdir=scratch-local</code>	Data are copied from the job input directory to the working directory on the computing node. The working directory is created at the beginning of the job by the batch system. When the job is completed, all data from the working directory is copied back to the job input directory. Eventually, the working directory will be deleted if the data transfer was successful.



# Data synchronization, cont.

## Suitable for analysis

Source	Meaning
<code>workdir=jobdir</code>	Job data are on shared storage.



# Running applications

# Request/use of resources

Batch system

Job

## Native batch system (PBSPro)

- **user specifies** required computing resources
- **user must ensure** that the job uses the assigned computing resources

## Infinity

- **user specifies** required computing resources
- **Infinity environment will ensure** correct starting of the job (selected applications only)
- (other tasks) **user must ensure** that the job uses the assigned computing resources

# Gaussian

The **gaussian** package contains tools for quantum chemical calculations. Detailed description can be found on <http://www.gaussian.com>



# gaussian, autodetection

Infinity can recognize the gaussian job type. The job script is automatically created and the input file is automatically updated according to requested resources.

```
$ module add gaussian  
$ psubmit default input.com ncpus=4
```



gaussian input file (must have .com extension), **this is NOT a job script!**

## Autodetection:

- job script is created automatically with correct gaussian binary name (g98, g03, g09, g16)
- %NProcShared is added or updated in the input file
- check if only single node is requested (parallel execution is limited to a single node)

```
[kulhanek@wolf 01.opt]$ psubmit default opt.com
```

```
Job name      : opt  
Job ID       : 34162.wolf-pbs.ncbr.muni.cz  
Job title    : opt (Job type: gaussian [gaussian:16.C1])  
Job input dir : wolf15.ncbr.muni.cz:/home/kulhanek/C7800/ProjectA/01.monomer/01.opt  
Job key      : 0fc3529f-52e8-fd61-5afd-fc9db76759d2  
Job project  : -none- (Collection: -none-)
```

# gaussian – single/parallel execution

The only difference between sequential and parallel execution is in the resource specification during psubmit. **The input data are the same!**

```
$ psubmit default opt.com ncpus=1
```

↑  
it can be omitted

Computational node:

S	%CPU	%MEM	TIME+	COMMAND
R	100	1.2	1:01.25	l502.exe
S	0	0.1	1:38.57	pbs_mom

```
$ psubmit default opt.com ncpus=4
```

Computational node:

S	%CPU	%MEM	TIME+	COMMAND
R	399	1.1	0:49.38	l502.exe
S	0	0.1	0:00.90	init

# gaussian, manual script preparation

```
#!/bin/bash

# activate gaussian module
module add gaussian:16.C1

# execute g16
g16 input
```

input file **input.com** must contain specification for number of CPUs requested for parallel execution (this number MUST be consistent with resource specification via **psubmit** command).

```
%NProcShared=4
```

```
$ psubmit default test_gaussian ncpus=4
```

**NOT RECOMMENDED**

Possible solution: psanitize command

<https://infinity.ncbr.muni.cz/whitezone/isoftrepo/fcgi-bin/isoftrepo.fcgi?action=module&site=cmng&module=gaussian>