

Introduction to Condor

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<http://www.cs.wisc.edu/condor>



Condor
High Throughput Computing

What is Condor?

- Condor is a system designed for "High Throughput Computing".
- It manages large numbers of jobs for users on lots of machines automatically. Especially useful for running lots of jobs over a long period of time.
- Developed and supported by University of Wisconsin (> 30 full time faculty and staff). Over 300 pools worldwide running on over 8500 CPUs.



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Some advantages of Condor:

- Balances requests for people, groups, and machines dynamically.
- You can manage a very large set of similar jobs with just one file. Condor runs them all sends you email when done etc.
- If job is linked specially, jobs can restart after computer crashes etc. at the exact point they left off, also they can move among computers and even run on computers where you don't have an account.
- Remote Condor pools can be linked together.



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How to run a job (A simple submit file):

```
# condor job control file
Executable    = my_prog
universe      = vanilla
getenv        = true

log            = condor-out.log

Arguments     = file1.dat

Output        = output.log
Error         = output.log

Queue 1
```

Use command: `condor_submit condor_job`



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How to process many files in one job:

condor job control file

Executable = my_prog

universe = vanilla

getenv = true

log = condor-out.log

Output = output-01.log

Error = output-01.log

Arguments = run-01.dat

Queue 1

Output = output-02.log

Error = output-02.log

Arguments = file-02.dat

Queue 1



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How to use macro variables:

condor job control file

Executable = my_prog

universe = vanilla

getenv = true

log = condor-out.log

Output = output-\$(fileNumber).log

Error = output-\$(fileNumber).log

Arguments = file-\$(fileNumber).dat

fileNumber = 01

Queue 1

fileNumber = 02

Queue 1



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Sometime you can use \$(Process):

condor job control file

Executable = my_prog

universe = vanilla

getenv = true

log = condor-out.log

Output = output-\$(Process).log

Error = output-\$(Process).log

Arguments = file-\$(Process).dat

Queue 100

Please look at `condor_submit` manual for many more commands.



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Managing your jobs

- `condor_submit` (submit your job)
- `condor_q` (status of all your jobs)
- `condor_q -r` (status of running jobs [what machines])
- `condor_status` (What machines are available/busy)
- `condor_rm` (Remove a job)
- `condor_history` (Show the history of a previous job)



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Clusters and Processes

```
> condor_q -r
```

```
-- Submitter: sukap02.suketto.icrr.u-tokyo.ac.jp : <10.10.10.102:62501> :  
sukap02.suketto.icrr.u-tokyo.ac.jp
```

ID	OWNER	SUBMITTED	RUN_TIME	HOST(S)
217.0	walter	4/18 19:18	0+00:00:11	sukap58
217.1	walter	4/18 19:18	0+00:00:09	sukap94
217.2	walter	4/18 19:18	0+00:00:07	sukap58

```
>
```



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Looking at the history of the jobs

```
> condor_history 190  
(ClusterId == 190)
```

ID	OWNER	SUBMITTED	CPU_USAGE	ST	COMPLETED	CMD
190.1	walter	4/14 17:56	0+00:56:08	C	4/14 18:52	/home/walter/k2
190.0	walter	4/14 17:56	0+01:37:36	C	4/14 19:34	/home/walter/k2
190.2	walter	4/14 17:56	0+03:45:45	C	4/14 21:43	/home/walter/k2

```
>
```



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Our Condor Pool

- 200 CPUs (sukap00-99)
- Interactive processes/Condor on 00-09
- Condor Only on sukap10-99
 - atmpd priority on 40-59
 - lowe priority on 60-79
 - upmu priority on 80-99



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Where to get more info

<http://www-sk/~condor> (local pool status)

For more info and documentation look at the main Condor Site:

<http://www.cs.wisc.edu/condor>

<http://www.cs.wisc.edu/condor/quick-start.html>

<http://www.cs.wisc.edu/condor/manual/v6.2/>

(users manual and man pages)

The Condor team has been very helpful to us. We have made a special agreement so that Fujitsu can access the source code and be treated as an academic group so they can support our system.



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