#### Introduction to HTCondor

How to distribute your compute tasks and get results with high performance, keeping machines and site admins joyful

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28<sup>th</sup> August, 2019

### Overview

- Introduction
- 2 How HTCondor works and how it can be used
- What might go wrong...
- Hands-on tutorial!

Find this talk and the actual tutorial at:

https://git.io/gridka-2019-htcondor

### Welcome!

#### About me

- studied physics in Bonn, starting in 2007
- PhD finished in 2017 at the BGO-OD experiment located at ELSA in Bonn (Hadron Physics, photoproduction)
   Focus on software development (C++ / ROOT)
- since 2017: IT dep. of Physikalisches Institut at Uni Bonn
  - Central services (desktops, printers, web, virtualization...)
  - Grid-enabled computing cluster:
     used by HEP, theory, detector dev., photonics,...
     HTCondor & Singularity containers, CephFS, CVMFS,...
  - Automation of all services and machine deployments
  - Support for users
  - IT security

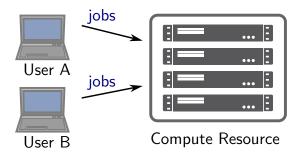
TL;DR: Feel free to ask both from user and admin point of view!



### **HTCondor**

- Workload Management system for dedicated resources, idle desktops, cloud resources, . . .
- Project exists since 1988 (named Condor until 2012)
- Open Source, developed at UW-Madison, Center for High Throughput Computing
- Key concepts:
  - 'Submit Locally. Run globally.' (Miron Livny)
    One interface to any available resource.
  - Integrated mechanisms for file transfer to / from the job
  - 'Class Ads', for submitters, jobs, resources, daemons, . . . Extensible lists of attributes (expressions) — more later!
  - Supports Linux, Windows and MacOS X and has a very diverse user base
    - CERN community, Dreamworks and Disney, NASA,...

## What is a workload manager?



(e.g. local cluster, desktops, cloud)

- takes care of collecting user's requirements
- prioritization / fair share
- enforcing limits
- collect resource information
- distribute jobs efficiently
- monitor status for users and admins



## Why HTCondor?

### **High Throughput Computing**

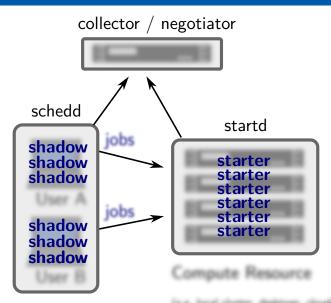
many jobs, usually loosely coupled or independent, goal is large throughput of jobs and / or data

#### High Performance Computing

tightly coupled parallel jobs which may span several nodes and often need low-latency interconnects

- HTCondor can do both (HPC-like tasks need some 'tuning')
- HPC community: Slurm (less flexible, but easier to get up and running for HPC!)
- ⇒ Let's have a look at how HTCondor works.

### **Structure of HTCondor**



### **HTCondor's ClassAds**

- Any submitter, job, resource, daemon has a ClassAd
- ClassAds are basically just expressions (key = value)
- Dynamic evaluation and merging possible

#### Job ClassAd

```
Executable = some-script.sh
+ContainerOS="CentOS7"

Request_cpus = 2
Request_memory = 2 GB
```

Request\_disk = 100 MB

#### Machine ClassAd

```
Activity = "Idle"
Arch = "X86 64"
Cpus = 8
DetectedMemory = 7820
Disk = 35773376
has avx = true
has_sse4_1 = true
has sse4 2 = true
has_ssse3 = true
KFlops = 1225161
Name = "slot1@htcondor-wn-7"
OpSys = "LINUX"
OpSysAndVer = "CentOS7"
OpSysLegacy = "LINUX"
Start = true
State = "Unclaimed"
                      UNIVERSITÄT BONN
```

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### **HTCondor's ClassAds**

- Job and Machine ClassAd extended / modified by HTCondor configuration
- Merging these ClassAds determines if job can run on machine
- Examples for dynamic parameters:
  - Select a different binary depending on OS / architecture
  - Machine may only want to 'Start' jobs from some users
- You can always check out the ClassAds manually to extract all information (use the argument -long to commands!)
- To extract specific information, you can tabulate specific attributes:

```
$ condor_q -all -global -af:hj Cmd ResidentSetSize_RAW

→ RequestMemory RequestCPUs

ID Cmd ResidentSetSize_RAW RequestMemory RequestCPUs

2.0 /bin/sleep 91168 2048 1
```

## What HTCondor needs from you...

#### A job description / Job ClassAd

Resource request, environment, executable, number of jobs,...

```
Executable = some-script.sh
Arguments = some Arguments for our program $(ClusterId) $(Process)
Universe = vanilla
Transfer_executable
                        = True
Error
                        = logs/err.$(ClusterId).$(Process)
                        = input/in.$(ClusterId).$(Process)
#Input
                        = logs/out.$(ClusterId).$(Process)
Output
                        = logs/log.$(ClusterId).$(Process)
Log
+ContainerOS="CentOS7"
Request_cpus = 2
Request_memory = 2 GB
Request_disk = 100 MB
Queue
```

#### some-script.sh

- Often, you want to use a wrapper around complex software
- This wrapper could be a shell script, python script etc.
- It should take care of:
  - Argument handling
  - Environment setup (if needed)
  - Exit status check (bash: consider -e)
  - Data handling (e.g. move output to shared file system)

```
#!/bin/bash
source /etc/profile
set -e
SCENE=$1

cd ${SCENE}
povray +V render.ini
mv ${SCENE}.png ...
```

### Submitting a job

```
$ condor_submit myjob.jdl
Submitting job(s)..
1 job(s) submitted to cluster 42.
```

There are many ways to check on the status of your job (we will try them in the tutorial):

- condor\_tail -f can follow along stdout / stderr (or any other file in the job sandbox
- condor\_q can access job status information (memory usage, CPU time,...)
- log file contains updates about resource usage, exit status etc.
- condor\_history provides information after the job is done
- condor\_ssh\_to\_job may allow to connect to the running job

### **Advanced JDL syntax**

```
Executable = /home/olifre/advanced/analysis.sh
Arguments = "-i '$(file)'"
Universe = vanilla
if $(Debugging)
    slice = [:1]
    Arguments = "$(Arguments) -v"
endif
Error = log/$Fn(file).stderr
Input = $(file)
Output = log/$Fn(file).stdout
Log = log/analysis.log
Queue FILE matching files $(slice) input/*.root
```

HTCondor also offers macros and can queue based on files. Can you guess what happens if you submit like this?

```
condor submit 'Debugging=true' analysis.jdl
```



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### **DAGs: Directed Acyclic Graphs**

- Often, jobs of different type of an analysis chain depend on each other
   Example: Monte Carlo, comparison to real data, Histogram merging,...
- These dependencies can be described with a DAG
- Condor runs a special 'DAGMAN' job which takes care of submitting jobs for each 'node' of the DAG, check status, limit idle and running jobs, report status etc. (like a Babysitter job)
- DAGMAN comes with separate logfiles, DAGs can be stopped and resumed

We will see an example in the tutorial!



### **Problems and inefficiencies**

- Theoretically, users should not need to care about cluster details...
- Jobs could transfer all their data with them, and back but this does not scale for GB of data, thousands of files for thousands of jobs
- Jobs need to take care to be 'mobile' and run in the correct environment

### Some setup details can not be ignored for efficient usage

Let's have a short look at elements of computing clusters and how (not) to design your jobs!

## A typical HTC cluster: I/O intensive loads

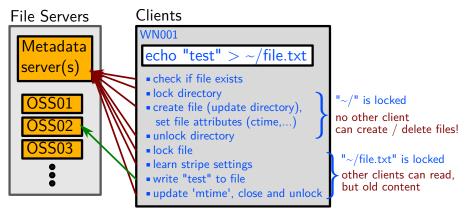
- Shared / parallel file system for data, job input and output CephFS, Lustre, BeeGFS, GPFS,...
- Often, also a second file system (e.g. to distribute software) CVMFS. NFS. . . .
- Usually, local scratch disks in all worker nodes 'classic' file system such as ext4
- Often, dedicated submit nodes, data transfer nodes etc.
- ⇒ Lots of differently behaving file systems!

### Working with a shared file system

#### Common sources of woes

- Excessive file metadata operations
  - Syscalls: open, close, stat, fsync...)
    use strace to diagnose and debug
- Storing or reading many small files from shared FS
   There is usually a dedicated place for software (more later).
- Destructive interference between jobs
  - Opening an input file exclusively
  - Writing to the very same output file

### Working with a shared file system



- x number of running jobs,
- x number of metadata accesses

### Working with a shared file system

#### **Common solutions**

- Use a different file system for software (many small files!) *CVMFS, NFS,...*
- Most software is (likely) already provided by cluster admins —
  use it!
  - They know how to compile best for the available hardware.
- Do not install everything from scratch (e.g. pip install "everything")
- Package quickly changing software builds in a tarball, extract it to scratch disk in the job wrapper script Advantage: Consistent software state for all jobs.
- Have jobs write to scratch first and move to shared FS later
   Advantage: If job is evicted, no broken output file.

### Working with a workload manager

#### Common sources of woes

- Mismatched resource request and usage (more later)
- Hefty / bad use of condor file transfer, for example: Shared FS accessible from submit machine, transferring files from / to there
- Badly suited job runtimes
  - too short Overhead per job causes inefficiency, some workload managers overload easily too long Unless the job does checkpointing, very sensitive to any disturbance, operational issues (kernel updates / reboots etc.)
- Frameworks which create thousands of JDL files and wrapper scripts (instead of using flexible syntax or Python API)

### Working with different environments

### How to compile code?

- Some resources may only be available via interactive jobs
  - Advantage for admins: No separate bare metal machines
  - Advantage for you: Environment the same as in the job!
- Compile the code, pack it into a tarball, send to shared FS / condor file transfer
- Can be automated with scripts / if offered, job start hooks (like '.bashrc')

### Advantages of this approach

- Portable and stable job executables
- If combined with containers and 'mobile data': Mostly cluster independent jobs possible

### Mismatched resource requests

### Mismatched CPU request

Often caused by software using all 'visible' cores — configure!

```
export NUMEXPR_NUM_THREADS=1
export MKL NUM THREADS=1
export OMP_NUM_THREADS=1
```

- Admins may export these variables for you...
- Too many threads: Congestion, may affect other jobs

#### Mismatched memory request

- Depending on configuration, may lead to swapping ⇒ hefty slowdown (affects also other jobs)
- Swap usage not visible in HTCondor Ads (yet)
- Admins could also set a hard limit (no swap) ⇒ job killed

### What about other resources?

### Disk Space

- Disk space is not 'consumable' in HTCondor
- Usually, this affects scratch space only (job working directory)
- Commonly, not an actual issue (shared file systems have quotas)
- More common is disk overload due to heavy syscalls / many small files / swap

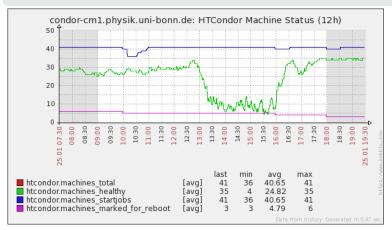
#### **CPU** cache thrashing

Commonly ignored issue — limiting e.g. CPU cache usage not supported by HTCondor yet (but there are plans)!

### Common tricks used by admins

#### Node health check

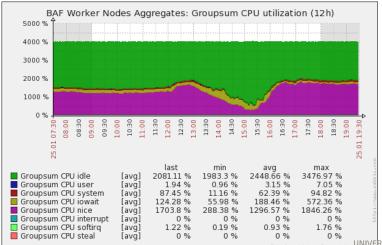
Detects unhealthy node from error or misbehaving jobs.



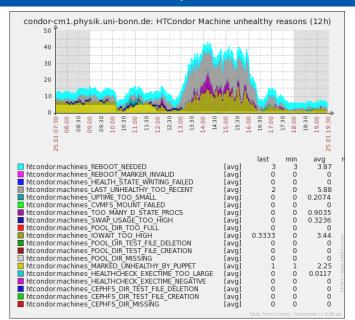
## Common tricks used by admins

#### Node health check

Fights against spread of inefficiencies / overload.



### Common tricks used by admins



### Conclusion

- HTCondor is very flexible you can check out configuration via ClassAds!
- Each cluster may be slightly different (CERN has job flavours to define job runtime, Bonn has containers with different environments,...)
- We will learn job submission today to run efficiently, you also need to know your software and the cluster

# Ask questions any time!

And now, get started at:

https://git.io/gridka-2019-htcondor



for your attention!

Thank you