# Introduction to HTCondor

**GPU Cluster** 

Faculty MI

#### Overview Infrastructure





**2** x Intel Xeon E5-2698 v4 20 cores [2.2 GHz]: **40 cores** 

**2** x Intel Xeon E5-2690 v4 28 cores [2.6 GHz]: **56 cores** 

**2** x Intel Xeon E5-2698 v4 20 cores [2.2 GHz]: **80 cores** 

**2** x Intel Xeon E5-2698 v4 20 cores [2.2 GHz]: **80 cores** 

**2** x AMD EPYC 7742 64 cores [2.2 GHz]: **256 cores** 







8 x Nvidia Pascal P100 PCle

Double-Precision 8 x 4.7 = 37.6 teraflops Single-Precision 8 x 9.3 = 74.4 teraflops Half-Precision 8 x 18.7 = 149.6 teraflops

8 x Nvidia Pascal P100 SXM

Double-Precision 16 x 5.3 = 84.8 teraflops Single-Precision 16 x 10.6 = 169.6 teraflops Half-Precision 16 x 21.2 = 339.2 teraflops

8 x Nvidia Pascal P100 SXM

Double-Precision  $16 \times 5.3 = 84.8$  teraflops Single-Precision  $16 \times 10.6 = 169.6$  teraflops Half-Precision  $16 \times 21.2 = 339.2$  teraflops

8 x Nvidia Volta V100 SXM

 $\begin{array}{lll} \mbox{Double-Precision } 16 \times 7.8 & = 124.8 \ \mbox{teraflops} \\ \mbox{Single-Precision } 16 \times 15.7 & = 249.6 \ \mbox{teraflops} \\ \mbox{Deep-Learning } 16 \times 125 & = 2000 \ \mbox{teraflops} \\ \end{array}$ 

8 x Nvidia Ampere SXM

 Double-Precision
 16 x 9.7
 = 155.2 teraflops

 Single-Precision
 16 x 19.5
 = 312 teraflops

 Deep-Learning
 16 x 312
 = 4992 teraflops

each x 512 GB

each x 768 GB

each x 512 GB

each x 512 GB

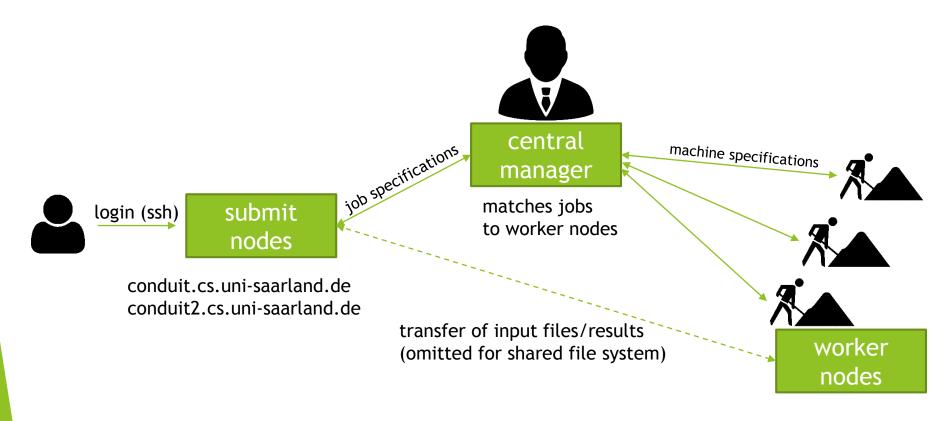
each x 1 TB

#### What is HTCondor?



- Software that runs and schedules tasks on computers
- Users can submit tasks to a queue from submit nodes
- ► HTCondor schedules them to run worker nodes
- Worker nodes can have different underlying architectures, e.g. different numbers and types of CPUs, availability of GPUs or not, different RAM size, etc.
- Users must define the requirements for their job (e.g. number of GPUs, needed RAM, ...) using a submit file

### Overview submit process



users of the GPU cluster can only login to the submit nodes

#### available CPU/GPU worker nodes

	GPU Type	# GPUs	CPU Type	# CPU Cores (incl. Hyper- Threading)	RAM
thor.cs.uni-saarland.de (DGX-1)	Tesla V100-SXM	8	Intel Xeon CPU E5- 2698 v4	80	512 GB
loki.cs.uni-saarland.de (DGX-1)	Tesla V100-SXM	8	Intel Xeon CPU E5- 2698 v4	80	512 GB
lofn.cs.uni-saarland.de	Tesla P100-SXM	8	Intel Xeon CPU E5- 2698 v4	80	512 GB
idun.cs.uni-saarland.de	Tesla P100-SXM	8	Intel Xeon CPU E5- 2698 v4	80	512 GB
uller.cs.uni-saarland.de	Tesla P100-PCIE	8	Intel Xeon CPU E5- 2698 v4	80	512 GB
tenos.cs.uni-saarland.de	Tesla P100-SXM	8	Intel Xeon CPU E5- 2690 v4	56	768 GB
thera.cs.uni-saarland.de	Tesla P100-SXM	7	Intel Xeon CPU E5- 2690 v4	56	768 GB
urd.cs.uni-saarland.de (DGX A100)	A-100-SXM	8	AMD EPYC 7742	256	1 TB
vidar.cs.uni-saarland.de (DGX A100)	A-100-SXM	8	AMD EPYC 7742	256	1 TB

#### Submitting a Job to the Cluster

- login on {conduit, conduit2}.cs.uni-saarland.de with your SIC LDAP credentials using ssh (only reachable from within the university network!)
- you will find a folder condor\_tutorial with some example submit files in your home directory
- ▶ /home is an NFS mount and shared between all nodes in the cluster
- this means that large files don't have to be copied between submit and worker nodes several times
- we only allow execution of jobs that use the HTCondor docker universe
- when you submit your job, condor will check if a worker node can be matched to run your job
- your job will stay idle as long as there is no worker available for running the job
- your job will be held if something obvious (e.g. missing input files) would prevent it from running successfully

#### Submitting a Job to the Cluster

- if your job enters the state running, condor creates a job specific scratch directory on the worker node that runs your job
- ▶ the scratch directory is mounted in the docker container that runs your job
- you can also prompt condor to mount the /home NFS mount into the container with a special flag in the submit file
- Jobs in the docker container are run as your user so that you have access to the files in the NFS share with your UID/GID
- this also means that you have no superuser privileges
- ▶ after the job is completed, condor will automatically clean up (destroy the docker container, remove the scratch directory)
- result files are either transferred back to the directory from where you submitted the job or when using the shared filesystem will be in your home where you directed the output

#### Example Submit File

file: \$HOME/condor\_tutorial/tf\_matmul\_docker.sub

```
universe
                         = docker
docker_image _____
                         = tensorflow/tensorflow:latest-gpu
executable
                         = tf matmul docker.py
output
                         = tf_matmul.$(ClusterId).$(ProcId).out
                         = tf_matmul.$(ClusterId).$(ProcId).err
error
                         = tf_matmul.$(ClusterId).log
log
should_transfer_files
                         = YES
when_to_transfer_output = ON_EXIT
request_GPUs = 2
request_CPUs = 1
request_memory = 1G
requirements = UidDomain == "cs.uni-saarland.de"
                                                     needed for mounting
                                                      /home in docker container
+WantGPUHomeMounted = true
queue 10
            puts this job 10x in the queue
            default is queue or queue 1 to run this job one time
```

#### Submit File

- universe must be "docker" or your job will never be run on any worker node
- you can use any available docker image that are freely accessible from docker hub, nvidia, etc.
- executable is the script file that will be run upon starting the container
  - ▶ if it is a relative path (as in the example) it will be copied by condor to the scratch directory and mounted in the docker container
  - ▶ if it is an absolute path, condor is taking it as a path to a file within the container
  - can be left out/commented out if your image defines an entrypoint that runs by default
- output, error, log will contain the messages from stdout, stderr, and the condor log messages for your job
- submit your job to the queue:
  condor\_submit <my\_submit\_file.sub>

### The condor queue

- condor\_q allows you to watch the state of your jobs
- idle jobs wait for free resources

## The condor queue

if a job is put into hold, something was wrong: check it with condor\_q -hold <jobid>

# The condor queue

```
The Requirements expression for job 360.004 is
   (UidDomain == "cs.uni-saarland.de") && TARGET.HasDocker && (TARGET.Disk >= RequestDisk)
&& (TARGET.Memory >= RequestMemory) && (TARGET.GPUs >= RequestGPUs) &&
(TARGET.HasFileTransfer)
Job 360.004 defines the following attributes:
   DiskUsage = 1
   RequestDisk = DiskUsage
   RequestGPUs = 2
   RequestMemory = 1024
The Requirements expression for job 360.004 reduces to these conditions:
      Matched Condition
             9 UidDomain == "cs.uni-saarland.de"
             7 TARGET.GPUs >= RequestGPUs
             5 [0] && [7]
  0.004: Job is held.
Hold reason: Error from slot1_1@urd.cs.uni-saarland.de: Docker job has gone over memory
Last successful match: Mon Mar 1 16:21:58 2021
 60.004: Run analysis summary ignoring user priority. Of 11 machines,
     6 are rejected by your job's requirements
      O reject your job because of their own requirements
     0 match and are already running your jobs
```

get more detailed information about your job matching to available workers with:

```
condor_q -analyze
condor q -better
```

reason for holding the job

# condor\_qedit / condor\_hold / condor\_release

- adjust job requirements with condor qedit "on-the-fly"
- release held job: condor\_release <jobid>

```
[user@conduit2 condor_tutorial]$ condor_qedit 360 RequestMemory 2024
Set attribute "RequestMemory" for 2 matching jobs.
```

[user@conduit2 condor\_tutorial]\$ condor\_release 360
All jobs in cluster 360 have been released

- you can also put your jobs into hold with condor\_hold <jobid>
- when a job is released it gets put back into idle state and the matchmaking takes place again

### condor\_rm

if you need to delete your jobs from the queue, you can use:

```
condor_rm <jobid>  //only remove job with jobid
condor_rm -a  //delete all your jobs from the queue
```

# condor\_status - overview of claimed and idle worker nodes

Name	0pSys	Arch	State	Activity	LoadAv	Mem	ActvtyTime
slot1@loki.cs.uni-saarland.de			Unclaimed	_	0.000	515882	0+05:25:42
slot1@tenos.cs.uni-saarland.de		_	Unclaimed		0.000	770858	0+05:26:18
slot1_1@tenos.cs.uni-saarland.de			Claimed	Busy	0.000	1024	0+00:00:04
slot1_2@tenos.cs.uni-saarland.de			Claimed	Busy	0.000	1024	0+00:00:04
slot1_3@tenos.cs.uni-saarland.de	LINUX	X86_64	Claimed	Busy	0.000	1024	0+00:00:04
slot1@thera.cs.uni-saarland.de	LINUX	X86_64	Unclaimed	Idle	0.000	770858	<b>0+05:26:07</b>
slot1_1@thera.cs.uni-saarland.de	LINUX	X86_64	Claimed	Busy	0.000	1024	0+00:00:01
slot1_2@thera.cs.uni-saarland.de	LINUX	X86_64	Claimed	Busy	0.000	1024	0+00:00:01
slot1_3@thera.cs.uni-saarland.de	LINUX	X86_64	Claimed	Busy	0.000	1024	0+00:00:01
slot1@thor.cs.uni-saarland.de	LINUX	X86_64	Unclaimed	Idle	0.000	515882	0+05:25:51
interactive2@uller.cs.uni-saarland.de		X86_64	Unclaimed	Idle	0.000	103175	0+00:38:36
interactive3@uller.cs.uni-saarland.de		_	Unclaimed		0.000	103175	0+05:26:25
interactive4@uller.cs.uni-saarland.de			Unclaimed		0.000	103175	0+05:26:25
interactive5@uller.cs.uni-saarland.de		_	Unclaimed		0.000	103175	0+05:26:25
slot1@uller.cs.uni-saarland.de		_	Unclaimed		0.000	101127	0+05:26:25
slot1_1@uller.cs.uni-saarland.de			Claimed	Busy	0.000	1024	0+00:00:05
slot1_2@uller.cs.uni-saarland.de			Claimed	Busy	0.000	1024	0+00:00:05
slot1@urd.cs.uni-saarland.de		_	Unclaimed			1030857	0+05:18:46
slot1_1@urd.cs.uni-saarland.de		_	Claimed	Busy	0.000	1024	0+00:00:04
slot1@vidar.cs.uni-saarland.de		_			0.000	1030857	0+05:18:19
slot1_1@vidar.cs.uni-saarland.de	LINUX	X86_64	Claimed	Busy	0.000	1024	0+00:00:01

#### Interactive Jobs

- you can use interactive jobs to debug your code before submitting large/many jobs to the queue
- the number of parallel running interactive jobs are limited to 4
- only one of our worker nodes allows interactive jobs
- interactive jobs are killed automatically after one hour to allow other users to get an interactive slot
- same syntax as normal submit, just add '-i' as additional parameter
- number of GPUs/CPUs is restricted to 1 for an interactive slot
- Please note that trying to use a docker image with a predefined entrypoint (e.g. hello-world) might not work as interactive job (exits the container after the script runs)

### Further reading/tutorials

This was just a very superficial glimpse on the options that HTCondor provides. You can find more detailed information in the official HTCondor docs:

- https://htcondor.readthedocs.io/en/latest/
- Submitting a job: <a href="https://htcondor.readthedocs.io/en/latest/users-manual/submitting-a-job.html">https://htcondor.readthedocs.io/en/latest/users-manual/submitting-a-job.html</a>
- Managing a job: <a href="https://htcondor.readthedocs.io/en/latest/users-manual/managing-a-job.html">https://htcondor.readthedocs.io/en/latest/users-manual/managing-a-job.html</a>