

# Tier1 and HTCondor resources usage

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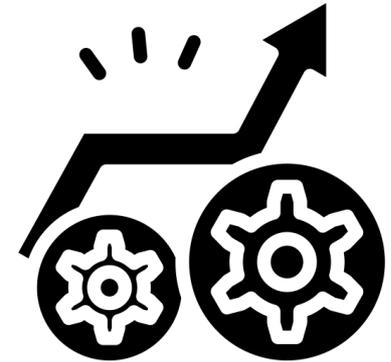
# Tier1

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From this year, we started moving all our files and resources to Tier1

- **Tier3 will not be our reference anymore**
- Tier1 is the main INFN computing service
- Login through bastion, two dedicated UIs for FOOT
  - ui01-foot (ui01-foot.cr.cnaf.infn.it)
  - ui02-foot (ui02-foot.cr.cnaf.infn.it)
- ROOT (6.26/08) already available (/opt/exp\_software/foot/root)
- SHOE installation is very straightforward
- Detailed instructions on environment setup available on the [software page of the FOOT TWiki](#)



# Tier1 – Storage



Each folder on the Tier1 has a specific task

→ Optimized for large files

`/storage/gpfs_data/foot` : parent directory (~90 TB)  
`/storage/gpfs_data/foot/shared` : Experimental data  
`/storage/gpfs_data/foot/shared/SimulatedData` : MC simulation data  
`/storage/gpfs_data/foot/${USER}` : User data (create if not present!)

**DATA**

→ Optimized for small files

`/opt/exp_software/foot` : parent directory  
`/opt/exp_software/foot/${USER}` : User software (create if not present!)

**SW**

→ Optimized for small files

→ Daily backup but limited and shared by all the collaboration

`/home/FOOT-T3` : parent directory (~110 GB)  
`/home/FOOT-T3/${USER}` : User \$HOME

**HOME**

# Tier1 – Software

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The UIs serve only as **interfaces**

- Not designed for demanding tasks
- Main purpose is I/O
- Limited resources (not more than a laptop)  
8 cpu, 16 GB RAM
- Shared by all people working in the experiment

# Tier1 – Software



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- Not designed for demanding tasks
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**BE CAREFUL WHAT YOU LAUNCH  
ON THESE MACHINES!**

*(maybe just avoid launching stuff here if you can...)*



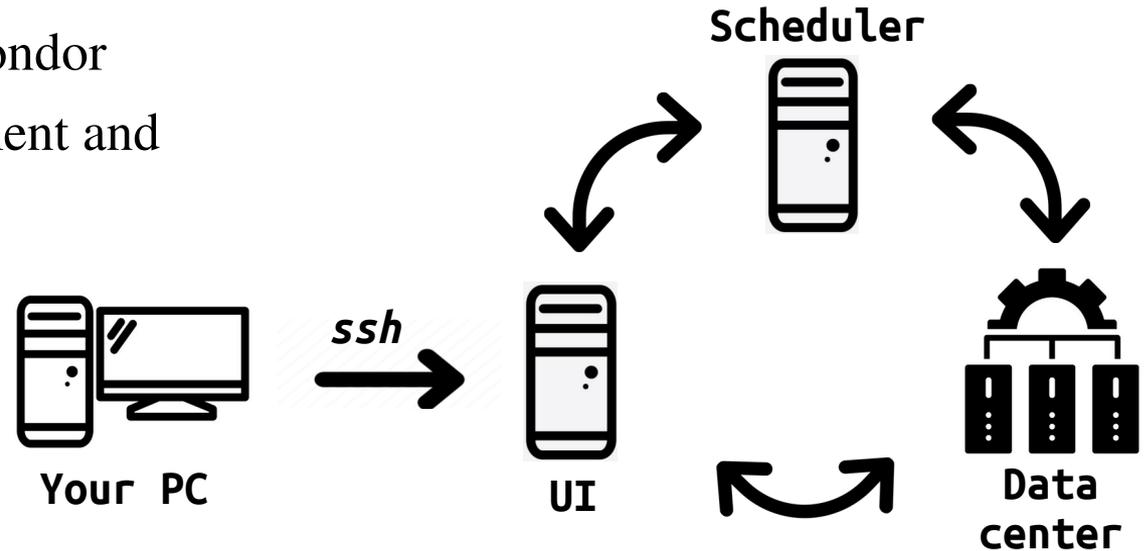
*So... How does one work on the Tier1?*

# Tier1 – HTCondor



The UIs are interfaced to the CNAF computing resources through the **HTCondor** batch system

- Best way to fully exploit the Tier1 resources
- From UI, a user can submit jobs to condor
- Job scheduler → handles the assignment and execution of the jobs
- What is needed to do that?
  - Executable file → command
  - Submit file → job parameters



**Scripts already available in SHOE to create aux files and submit jobs!**

# SHOE scripts – MC simulation processing



## shoe/Reconstruction/scripts/runShoeBatchT1\_MC.sh

### Arguments

- MC files processing
- Documentation inside the file
- **Can be run from any directory**
- **1 job = 20k events**
- **Campaign and run number are retrieved automatically!**
- Possible to process multiple files for a single run (-f option)
- **Mandatory to pre-set the FootGlobal.par and all configurations/calibrations!**  
(Maybe test DecodeGlb w/ ~100 evts before launching the script)

- i **Input file** (in /storage/gpfs\_data/foot/shared/SimulatedData)
- o **Output directory** (in /storage/gpfs\_data/foot/\${USER})
- m **Merge output files** (optional, default “0”)
- f **Use full statistics** (optional, default “0”)

### Example

```
./runShoeBatchT1_MC.sh -i /storage/gpfs_data/foot/shared/SimulatedData/12C_200_2023v2/12C_C_200_1_shoereg.root -o /storage/gpfs_data/foot/${USER}/results -m 1 -f 1
```

# SHOE scripts – Experimental data processing



## shoe/Reconstruction/scripts/runShoeBatchT1.sh

### Arguments

- Experimental data processing
- Documentation inside the file
- **Can be run from any directory**
- **1 job = 1 subfile**
- Possible to process multiple runs (even all!!) for a single campaign
- **Mandatory to pre-set the FootGlobal.par and all configurations/calibrations**  
(Maybe test DecodeGlb w/ ~100 evts before launching the script)

- i** Input directory (in /storage/gpfs\_data/foot/shared
- o** Output directory (in /storage/gpfs\_data/foot/\${USER})
- c** Campaign name
- r** First run number
- l** Last run number (optional)
- m** Merge output files (optional, default “0”)

### Example

```
./runShoeBatchT1.sh -i /storage/gpfs_data/foot/shared/DataGSI2021sync/ -o /storage/gpfs_data/foot/${USER}/results  
-c GSI2021 -r 4306 -l 4310 -m 1
```

# HTCondor jobs monitoring and handling



Jobs can be monitored through the “condor\_q” command:

```
condor_q -name sn-02
```

```
-- Schedd: sn-02.cr.cnaf.infn.it : <131.154.192.42:9618?... @ 11/27/23 17:23:48
OWNER          BATCH_NAME      SUBMITTED   DONE    RUN    IDLE  TOTAL JOB_IDS
zarrellafoott3 ID: 9934116    11/27 17:22   -      90     -     122 9934116.0-121
zarrellafoott3 ID: 9934118    11/27 17:22   -      -      1      1 9934118.0
zarrellafoott3 ID: 9934119    11/27 17:22   -      93     -     93 9934119.0-92
zarrellafoott3 ID: 9934120    11/27 17:22   -      -      1      1 9934120.0
zarrellafoott3 ID: 9934121    11/27 17:22   -     125     8    133 9934121.0-132
zarrellafoott3 ID: 9934123    11/27 17:22   -      -      1      1 9934123.0

Total for query: 351 jobs; 32 completed, 0 removed, 11 idle, 308 running, 0 held, 0 suspended
Total for zarrellafoott3: 351 jobs; 32 completed, 0 removed, 11 idle, 308 running, 0 held, 0 suspended
Total for all users: 29350 jobs; 27389 completed, 0 removed, 305 idle, 998 running, 658 held, 0 suspended
```

```
condor_q -name sn-02 -nobatch
```



Do not group jobs on same cluster

```
condor_q -name sn-02 -run
```



Display only running jobs with their runtime

You can check the output of a single running job in real time via “condor\_tail”

```
condor_tail -name sn-02 -f jobId
```

```
jobId = 9934116.0
        9934116.3
        9934121.130
```

The “-f” option lets you follow the output, you can omit it if you want a single printout

# HTCondor jobs monitoring and handling



Once all your jobs are completed, you **need** to remove them from the queue

```
condor_rm -name sn-02 $USER
```

```
condor_rm -name sn-02 -all
```

The “-name sn-02” can be omitted by adding the following line to your ~/.bash\_profile

```
export _condor_SCHEDD_HOST=sn-02.cr.cnaf.infn.it
```

If for some reason your jobs run for too long (>~1 hour), it is likely that something broke

- ☀ Usually SHOE’s “fault” → check if output files are created + condor\_tail
- ☀ Some jobs **will** go on “hold” after 2-3h → intended to free resources
- ☀ Other possible issues can be pointed out in the auxiliary files of each job (.out/.err/.log)

```
condor_transfer_data -name sn-02 -all
```

Further info and commands in the [CNAF-Tier1](#) and the [HTCondor](#) documentations

# HTCondor jobs monitoring and handling



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```
condor_transfer_data -name sn-02 -all
```

*If you can't find out why it is failing, try harder! And then obviously ask me if you need!*

# A possibility for production (finally!)



We now have the tools to process entire campaigns!

```
./runShoeBatchT1.sh -i /storage/gpfs_data/foot/shared/DataCNA02023sync/ -o  
/storage/gpfs_data/foot/${USER}/results -c CNA02023 -r 5957 -l 6313 -m 1
```



**Process CNA02023!**

Need to agree on a set of reconstruction parameters → **production!**

*(default in SHOE repo?)*

## I/O

```
EnableTree: y  
EnableFlatTree: n  
EnableHisto: y  
EnableTracking: y  
  
EnableSaveHits: y  
EnableRootObject: y  
EnableRegionMc: y  
EnableElecNoiseMc: ?
```

## Global Reco

```
EnableKalman: ?  
Kalman preselection strategy: ?  
N measure in global tracking: ?  
  
IncludeTOE: ?  
TOE cuts: ?  
  
IncludeStraight: ?  
Parameters: ?
```

## Detectors

```
IncludeDI: y  
IncludeST: y  
IncludeBM: y  
IncludeTG: y  
IncludeVT: y  
IncludeIT: y  
IncludeMSD: y  
IncludeTW: y  
IncludeCA: y
```

# Conclusions

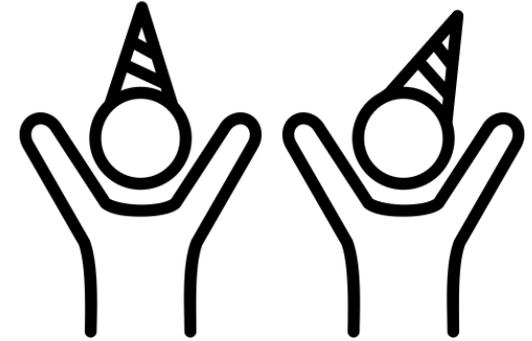
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Migrated to Tier1 resources

- ✓ SHOE installation straightforward
- ✓ Scripts for data and MC simulation processing w/ HTCondor available in SHOE
- ✓ Possibility now to go in production

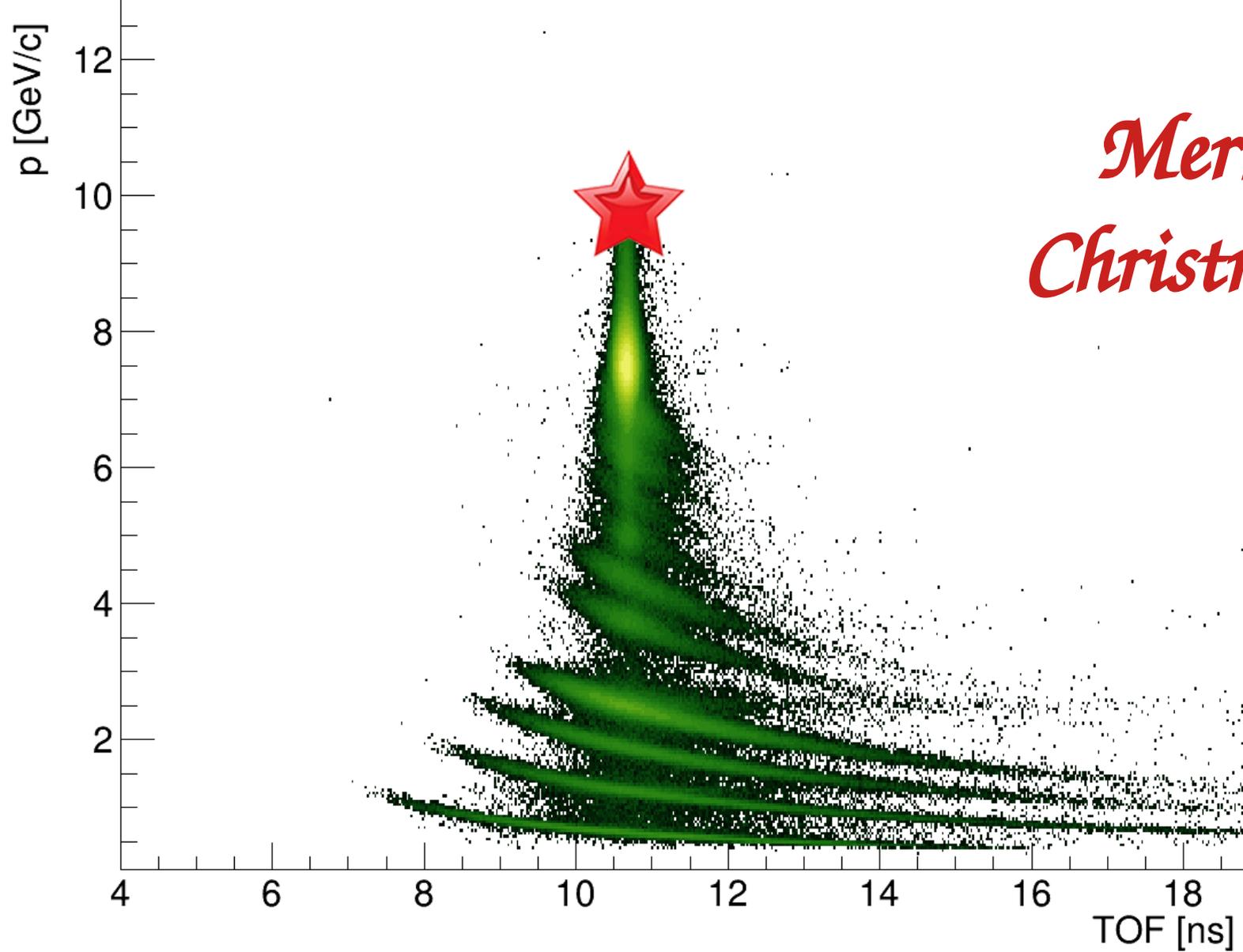
- Process everything w/ SHOE master branch
- Make processed files available for analysis
- Need for config/calib files ASA(R)P!



💡 Script for running macros on condor is in the works

💡 Need to decide workflow for MC input files production (FLUKA on Tier1?)

**Anyone who wants to contribute is welcome!!**



*Merry  
Christmas!*