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# German NAF: CMS user experience

#### Johannes Lange

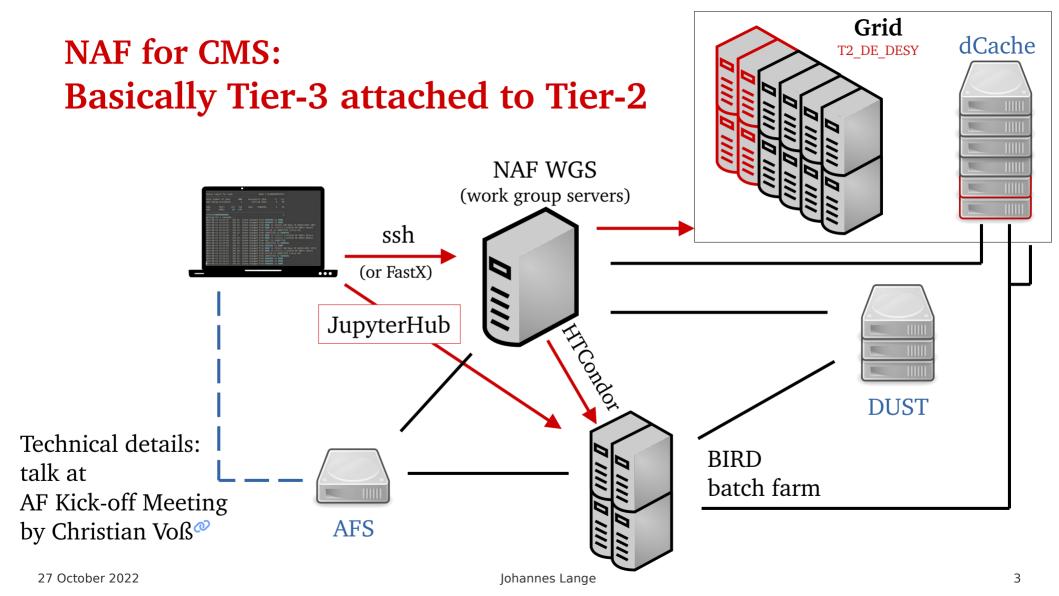
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## NAF: National Analysis Facility @DESY Hamburg

- Supposed to be used by the German HEP community
  - ATLAS, CMS, LHCb, ILC, CALICE, BELLE, HERA, smaller experiments
- CMS groups:
  - RWTH Aachen
  - DESY
  - Universität Hamburg
  - KIT





## NAF user storage



- AFS
  - /afs/desy.de/user/<u>/<user>
  - source code, documents etc.



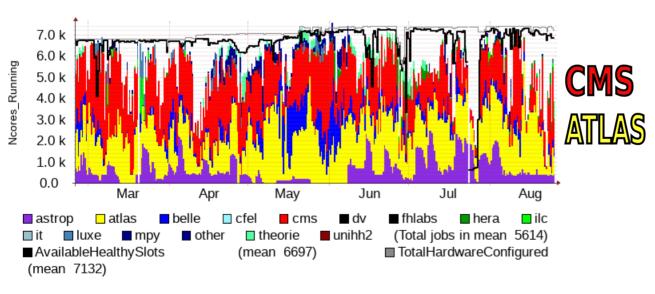
- DUST
  - /nfs/dust/cms/user/<user>
  - for large files (e.g. n-tuples),
     not intended for source code / many small files
- dCache
  - read-only NFS mount (WGS and worker nodes)
    /pnfs/desy.de/cms/tier2/store/[data|mc|user]
  - for large files (e.g. private productions, n-tuples)

27 October 2022

## NAF batch system: BIRD

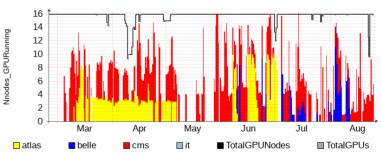
#### (Batch Infrastructure Resource at DESY)

- HTCondor with fair share between experiments
  - migrated from SGE 2017 2018
- largest number of HEP groups in Germany: ATLAS



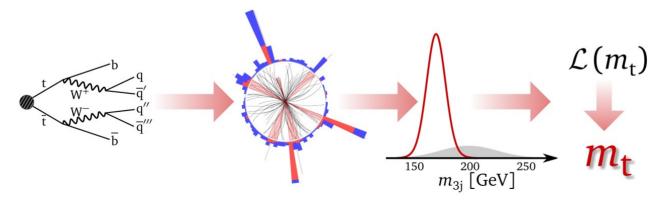
- Uni Hamburg CMS group integrates their resources in NAF
   → approx. equal share between ATLAS and CMS
- oversubscription for "lite" jobs (1 core, 2 GB mem, 3 hours runtime): more than fair share can be used, if resources are available

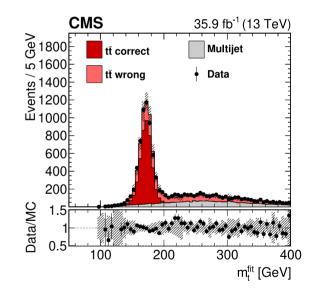
#### GPU-nodes available:



## Example analysis SM precision measurement: top quark mass, all-jets channel

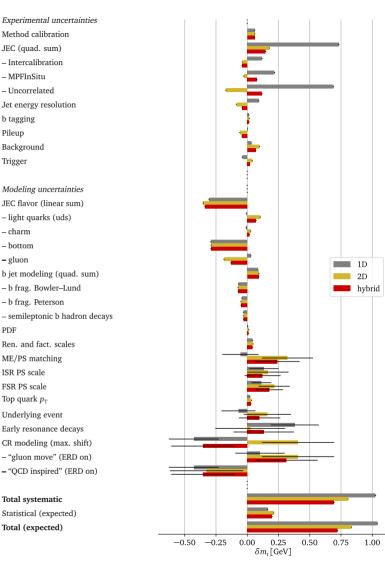
- completely performed @NAF, no grid-jobs
- computationally challenging:
  - huge QCD multijet background
  - combinatorics, kinematic fitting
  - systematic uncertainties: many variations





# Example analysis **Overview of tasks**

- Starting from MINIAOD data format (signal MC, background MC, data, background estimation from data)
- writing very specialized n-tuples (skimmed and slimmed) used for
  - plotting
  - statistical framework
- O(200) variations to process
- O(1000) pseudo-experiments per variation
- background estimation studies: different n-tuple format (slimmed)





# Example analysis **job submission: grid-control**<sup>@</sup>

- we advertise grid-control as standard job submission tool
- take care of job submission, checking and possibly resubmission
- great parameterization mechanism
- a number of batch systems are supported as backend

[arXiv:1707.03198]<sup>@</sup>

Status report for task:				demo / GC4884d5070fa7
	mber of jobs ng processed			Successful jobs: 22 11% Failing jobs: 0 0%
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# Example analysis Workflow

- CMS-central datasets: transferred to DESY dCache (formerly PhEDEx, now rucio)
- CMSSW-jobs (requiring full software stack)
  - write custom n-tuples to DUST
  - for background studies, write (large) n-tuples to dCache
- own C++/Python programs: do not depend on full CMSSW-stack, use n-tuples
  - pseudo-experiments, statistical procedures
  - background studies
  - plotting etc. (partially on WGS)

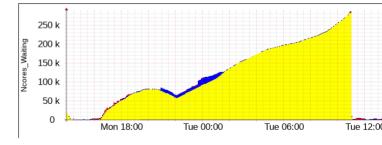
PhEDEx / Rucio WLC dCache DESY batch system dCache WGS user part DUS local dis

# Example analysis Some takeaways, pitfalls

- having data locally, mounted via NFS can be very convenient
  - (bachelor) students do not need CERN account, grid certificate
- for long-running CMSSW-jobs one should consider submitting to the grid (crab)
  - they "hurt" your priority in the batch system
     → more lightweight jobs down the line take longer to start
- sometimes users kill the HTCondor schedd by submitting too many jobs at the same time (becomes unresponsive with O(100k))
  - in grid-control easily controlled:
- we have different schedd nodes (1 ATLAS, 1 CMS, ...)
  - $\rightarrow$  not everybody is directly affected

in queue = 1000

[jobs]



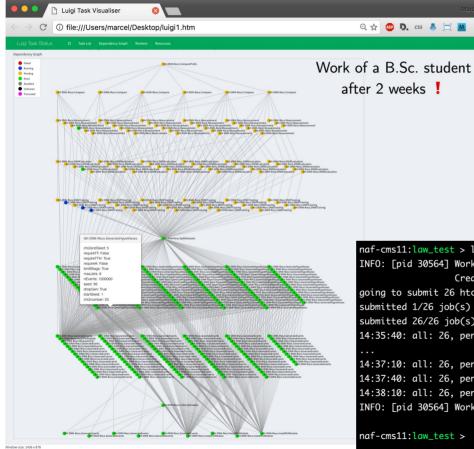
## JupyterHub @ NAF

Log in with DESY Account	Welcome to the JupyterHub for NAF User	ГS Г
Username: Password: Sign In	In order to login into the JupyterHub you must have your DESY credentials prepared for N Please follow the documentation of your experiment/group to gain full access to the NAF. You may also be interested in our other services, like the DESY supercomputer Maxwell.	Home Token jolange 🕞 Logout
– notebook ir	Select GPU node  Note: The <i>nafgpu</i> resource is needed for GPU nodes Jupyter Launch Modus Classical Notebook	
<ul> <li>useful for o</li> </ul>	Job Requirements e.g. Machine == 'batch1074.de Extra notebook CLI arguments e.gdebug	
— esp. tru	Environment variables (one per line)           YOURNAME=jolange	

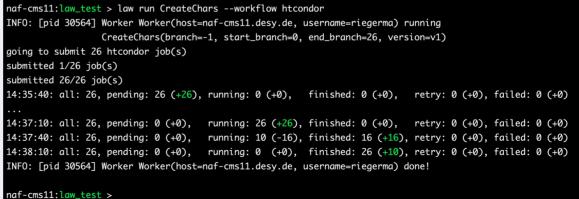
#### Material provided by Marcel Rieger

## Going more complex: law<sup>@</sup>

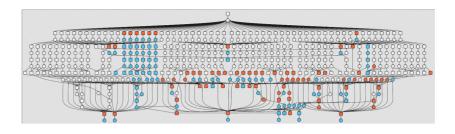




- based on *luigi* pipelining framework
- used by an increasing number of groups
- using HTCondor backend @NAF

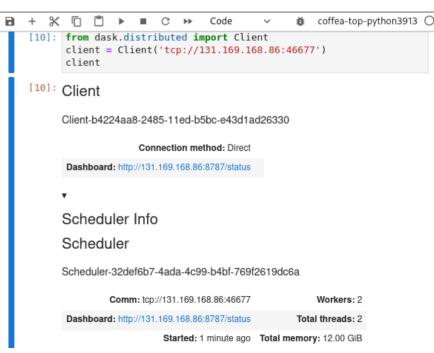


## dask-jobqueue





- attempt to make dask-jobqueue conveniently usable @NAF
- more interactive support of columnar analysis workflows
- spawn workers in existing HTCondor infrastructure
- very first steps
  - works on WGS with venv, conda, mamba, ...
  - can connect to started client from JupyterHub@NAF
- to be done
  - make it usable directly from JupyterHub@NAF (not configured for job-submission, yet)
  - monitor batch system usage to see if a special treatment (priority) for these jobs is needed



## **Summary**

- NAF is vital for German CMS analyzers
  - for many, grid jobs are not even necessary
- very different workflows possible
  - different analyses with different needs
  - different tools
- batch system is the work horse
  - backend to most tools

### Thanks for your attention!

