ctapipe, the GRID and I

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start off with some useful links (names are clickable)

DIRC Users Guide how to install dirac client, essential terminal commands and python methods

DIRAC JobMonitor for your jobs submitted by DIRAC; has very limited "resubmit" capabilities, can look at logs, stdout/stderr and output files on-the-fly

MC Prod3 Status description of which MC sets exist and how to obtain them

DIRAC from the command line

I assume you managed to set it up with the User Guide

first, define a bash function for convenience:

```
1 source-dirac (){
2    [[ "$PATH" =~ "dirac/pro/scripts" ]] || \
3         source ~/software/dirac/bashrc;
4         dirac-proxy-init
5    }
```

Retrieve a list of files of your GRID user and save it in a text file in the current directory: dirac-dms-user-lfns

upload a file to a specifiic storage element (SE):

dirac-dms-add-file /vo.cta.in2p3.fr/user/[initial]/[name]/test.txt test.txt CC-IN2P3-USER

download a file from the SE:

 $dirac\text{-}dms\text{-}get\text{-}file\ LFN:/vo.cta.in2p3.fr/user/[initial]/[name]/dir1/job.log$

remove a directory and all files within:

dirac-dms-clean-directory /vo.cta.in2p3.fr/user/[initial]/[name]/dir

create a replica on another SE:

dirac-dms-replicate-lfn/vo.cta.in2p3.fr/user/[initial]/[name]/test.txt DESY-ZN-Disk

A very convenient way to submit jobs to the GRID is through a python2 script. All the dirac-specific classes are made available by sourcing the dirac bashrc.

Note: This bashrc will set the python2 (!) version that ships with it as the default *python*, so only source it in a dedicated "submission terminal". It also messes up your *conda* upgrade process. I thought I messed up my installation a few times because of this...

My actual submit script is on my github.

submitting jobs to the GRID with DIRAC the preamble

- 1 from DIRAC. Core. Base import Script
- 2 Script.parseCommandLine()
- 3 from DIRAC.Interfaces.API.Dirac import Dirac
- 4 from DIRAC.Interfaces.API.Job import Job
- 5 dirac = Dirac()

sandbox uploads local files to the working directory of your GRID jobs

```
input sandbox = [
   # files from current directory; will be available
   # in working directory (WD) during GRID-job
8
9
    "helper_functions.py", "reconstruct.py",
10
11
   # file in a different directory; will be in WD (not in sub-dir.)
   "snippets/append_tables.py",
12
13
14
    # directory in current directory; will be in WD
15
   "modules",
16
17
   # some directory anywhere on disk; will be (you guessed it) in WD
   "/local/home/tmichael/software/jeremie_cta/"
18
    "sap-cta-data-pipeline/datapipe/",
19
20
21
   # file from a GRID SE (note the LFN prefix); will be in WD
22
   "LFN:/vo.cta.in2p3.fr/user/t/tmichael/cta/bin/mr filter/"
23
    "v3 1/mr filter"]
```

setting up a job object and submitting to the GRID

```
for run filelist in sliding window (filelist, window size = 10):
24
25
        i = lob()
26
        # runtime in seconds times 8 (CPU normalisation factor)
27
        j.setCPUTime(6 * 3600 * 8)
        # set a nice and unique name for your job
28
29
        j.setName(''.join([channel, mode, run token]))
30
31
        # store results on the SE; supports wildcards in filenames
32
        j.setOutputData(["output.log", "output *.h5"],
33
            # 'outputPath' relative path starting from your [name]
34
            outputSE=None, outputPath="output/path/")
35
36
        # files to upload to WD from current machine and other SEs
37
        j.setInputSandbox(input sandbox)
        . . .
54
        # submit the job to the GRID
        print('Submission_Result:_{}\n'.format(
55
              dirac.submit(j)['Value']))
56
```

submitting jobs to the GRID with DIRAC defining the executable

can be any command available in the WD, local scripts need to be added to the input sandbox to be uploaded

```
    38 # takes two strings, first is the executable to run,
    39 # second contains additional command line arguments
    40 j.setExecutable("reconstruct.py", command_line_arguments)
```

defining the input data we want to analyse

naive idea is to 'setInputData' and tell your executable to process them all:

```
38  j.setInputData(run_filelist)
39  j.setExecutable("reconstruct.py", "---input_*.simtel.gz")
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sends the job to the GRID node where files actually are stored

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- might be impossible if too many files; no SE has all the data
- running on only one/few files is not economical e.g. there are 45k prod3b-paranal-north-20deg-[gamma,proton] files, one file takes 10 min to run (too short), produces 45k 100 kB output files (takes three days only to download)

defining the input data we want to analyse

or add it to InputSandbox instead:

```
38  j.setInputSandbox(input_sandbox +
39      [input_file_on_SE_1 ,
40           input_file_on_SE_2 ,
41           ...])
42  j.setExecutable("reconstruct.py", "—input_*.simtel.gz")
```

sends your jobs anywhere and pulls data at the beginning to your WD

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- $\bullet \rightarrow$ if too many jobs try to download files in parallel, interface breaks
- \bullet \rightarrow jobs idle around for 3 h and then get killed
- local WD has limited size, download too much and you run out of space

defining the input data we want to analyse

what actually works for me

```
# leave input sandbox as is
38
39
   j.setInputSandbox(input sandbox)
   # loop over all files in the window
40
   for run file in run filelist:
41
42
        # wait for a randomly up to five minutes before starting
        sleep = random.randint(0, 5*60)
43
44
        j.setExecutable('sleep', str(sleep))
        # consecutively downloads the data files,
45
        j.setExecutable('dirac-dms-get-file', "LFN:"+run file)
46
47
        # ... processes them and deletes them again
        j.setExecutable("reconstruct.py",
48
            "--input" + basename(run_file))
49
        i.setExecutable('rm', basename(run_file))
50
   # merge the output files
51
   j.setExecutable("append_tables.py",
52
        "...".join(run_filelist)
53
```

other possibly useful things

```
# if there is a temporary problem at one site, switch it off
j. setBannedSites (['LCG. IN2P3-CC. fr'])
# your large-scale submission was interrupted?
# check which jobs are already there
running_ids = []
for status in ["Waiting", "Running"]:
    running ids += dirac.selectJobs(status=status,
                                    owner="tmichael")['Value']
running tokens = []
for id in running ids:
    # with many jobs, this could take a minute or two
    jobname = dirac.attributes(id)["Value"]["JobName"]
    running tokens.append(jobname)
# list of available storage elements:
"CC-IN2P3-USER", "DESY-ZN-USER", "CEA-USER", "LAPP-USER",
"CYF-STORM-USER", "CNAF-USER", "LPNHE-USER"
```