

# CRAB tutorial

## Cms Remote Analysis Builder

*Stefano Lacaprara, Federica Fanzago, Marco Corvo*

Department of Physics  
INFN and University of Padova

CRAB Tutorial, 4 February 2005



# Outline

## Intro

- What is this tutorial about
- Prerequisites
- Further information

## CRAB usage

- Getting CRAB
- Configuration
- Running CRAB
- Troubleshooting



# What is this tutorial about

and what is **not**

- ▶ Tutorial about CRAB Cms Remote Analysis Builder
  - ▶ A tool developed within Workload Management group open the **Grid** to the masses!
  - ▶ CRAB is aimed to give access to CMS analysts to all Data produced and available everywhere, using GRID (LCG) middleware
  - ▶ CRAB should hide as much as possible Grid complexities and subtleties to CMS user (if you could believe it ...)
- ▶ This is **not a tutorial about**
  - ▶ How to do analysis
  - ▶ How to use CMS software (ORCA)
  - ▶ general Grid usage



# What is this tutorial about

and what is **not**

- ▶ Tutorial about CRAB **C**ms **R**emote **A**nalysis **B**uilder
  - ▶ A tool developed within Workload Management group open the **Grid** to the masses!
  - ▶ CRAB is aimed to give access to CMS analysts to all Data produced and available everywhere, using GRID (LCG) middleware
  - ▶ CRAB should hide as much as possible Grid complexities and subtleties to CMS user (if you could believe it ...)
- ▶ This is **not a tutorial about**
  - ▶ How to do analysis
  - ▶ How to use CMS software (ORCA)
  - ▶ general Grid usage



# Prerequisites

- ▶ What you need to access the Grid and access CMS data:
- ▶ Work from an **User Interface** (UI)
- ▶ Have a valid Grid certificate
- ▶ Have a active grid-proxy on UI (grid-proxy-init)
- ▶ Your Virtual Organization (VO) must be (also) **CMS**
- ▶ Have CMS sw (ORCA) installed on UI
- ▶ **Know how to use ORCA interactively!**



## More information sources

- ▶ CRAB web page  
<http://cmsdoc.cern.ch/cms/ccs/wm/www/Crab>
- ▶ CRAB mailing list, for feedback and user support (you can register via SIMBA) [cms-wm-crab-feedback@cern.ch](mailto:cms-wm-crab-feedback@cern.ch)
- ▶ CRAB savannah web page: for bugs report and features request <https://savannah.cern.ch/projects/crab/>
- ▶ README (which comes with CRAB)
- ▶ on-line crab manual `-help`
- ▶ EIS support [support-eis@cern.ch](mailto:support-eis@cern.ch) for **GRID** related problems



# Get CRAB

## ► At CERN

- Any lxplus[7] node becomes an UI sourcing:  
`/afs/cern.ch/cms/LCG/LCG-2/UI/cms_ui_env.csh|sh`
- From any lxplus[7] node, just source:  
`$CMS_PATH/ccs/wm/script/Crab/crab.csh|sh`  
(it sources UI script is not done yet)
- Keep updated for you!

## ► Elsewhere

- Get it from CVS (working with a better distribution system...)
- `cmscvsroot CRAB`
- `cvs co -r <tag> UserTools`
- latest tag (04-Feb-2005) is CRAB\_0\_0\_7



## How to start

- ▶ From a UI (lxplus[7] is fine) write and test your code in the usual way, using an ORCA working area (`scram project ...`)
- ▶ Once you are happy, decide which remote dataset you want to access

- ▶ Which dataset are available?

- ▶ Look at CERN PubDB page

<http://cmsdoc.cern.ch/cms/production/www/PubDB/GetPublishedCollectionInfoFromRefDB.php>

- ▶ **Warning!!!** the data location service is ramping up: not all site are **today** up to date and accessible via CRAB
- ▶ Site **today** working fine: CNAF, LNL, BA; PIC almost; FNAL, IN2P3, CERN are working...
- ▶ LCG Worker nodes are RedHat7.3,  $\Rightarrow$  must use a RH7.3 UI: lxplus7 at CERN. Migration to SLC3 should take place soon





# CRAB configuration

- ▶ From user working area (e.g. `~/ORCA_8_7_1/src/Workspace`), issue the usual command `eval 'scram runtime -sh|csh'`
- ▶ Move to your Crab working area:
  - ▶ `UserTools/src` if CRAB via CVS
  - ▶ *not needed if at CERN and `crab.csh` sourced!*
- ▶ modify configuration file `crab.cfg`
- ▶ run `crab`



# CRAB configuration

- ▶ From user working area (e.g. `~/ORCA_8_7_1/src/Workspace`), issue the usual command `eval 'scram runtime -sh|csh'`
- ▶ Move to your Crab working area:
  - ▶ `UserTools/src` if CRAB via CVS
  - ▶ **not needed if at CERN and `crab.csh` sourced!**
- ▶ modify configuration file `crab.cfg`
- ▶ run `crab`



# CRAB configuration

- ▶ From user working area (e.g. `~/ORCA_8_7_1/src/Workspace`), issue the usual command `eval 'scram runtime -sh|csh'`
- ▶ Move to your Crab working area:
  - ▶ `UserTools/src` if CRAB via CVS
  - ▶ `not needed if at CERN and crab.csh sourced!`
- ▶ modify configuration file `crab.cfg`
- ▶ run `crab`



# CRAB configuration

- ▶ From user working area (e.g. `~/ORCA_8_7_1/src/Workspace`), issue the usual command `eval 'scram runtime -sh|csh'`
- ▶ Move to your Crab working area:
  - ▶ `UserTools/src` if CRAB via CVS
  - ▶ **not needed if at CERN and `crab.csh` sourced!**
- ▶ modify configuration file `crab.cfg`
- ▶ run `crab`



# CRAB configuration

- ▶ From user working area (e.g. `~/ORCA_8_7_1/src/Workspace`), issue the usual command `eval 'scram runtime -sh|csh'`
- ▶ Move to your Crab working area:
  - ▶ `UserTools/src` if CRAB via CVS
  - ▶ **not needed if at CERN and `crab.csh` sourced!**
- ▶ modify configuration file `crab.cfg`
- ▶ `run crab`



# CRAB configuration

- ▶ From user working area (e.g. `~/ORCA_8_7_1/src/Workspace`), issue the usual command `eval 'scram runtime -sh|csh'`
- ▶ Move to your Crab working area:
  - ▶ `UserTools/src` if CRAB via CVS
  - ▶ **not needed if at CERN and `crab.csh` sourced!**
- ▶ modify configuration file `crab.cfg`
- ▶ run `crab`



# crab.cfg

► Mandatory keys:

**dataset** The one you want to access

**owner** ditto

**executable** the one you used interactively! It is in your path since you did `eval 'scram runtime -csh'`, didn't you? If the executable uses some private libs, **crab** find them for you.

**orcarc\_file** the .orcarc file you used interactively: the very same! **crab** will change it for you

**total\_number\_of\_events** to be processed

**job\_number\_of\_events** number of event per job

**output\_file** your executable produces



# crab.cfg

► Might be useful:

`data_tiers` you want to access with your executable:  
“DST,Digi,Hit”. Require parents to be  
published in the same site with primary owner.  
Default is just primary data tier.

**PU not yet possible! WARNING:**  
**ExDSTStatistics access PU!!!**

`additional_input_files` to be send to WorkerNode (WN) with  
the jobs: comma (,) separated list

`ouput|log_dir` directory where crab will put output and log: if  
not set the default is  
`crab_0_<date>_<time>/res`





# Running CRAB

`crab.py -help` and read it!!!

`crab.py -create N` to create N jobs (no submission)

`crab.py -submit N -continue` to submit the job you have created  
(mind the `-continue`)

`crab.py -monitor -continue` to monitor (very primitive!) your jobs  
and get automatically the output retrieved when the  
jobs are finished

`crab.py -create all -submit all -monitor` to do all in just one  
command

If all is fine, present your work at meeting and offer some  
wine/beers/etc to crab team!

if problems, see next slides (and no beer for us...)



# Running CRAB

`crab.py -help` and read it!!!

`crab.py -create N` to create N jobs (no submission)

`crab.py -submit N -continue` to submit the job you have created  
(mind the `-continue`)

`crab.py -monitor -continue` to monitor (very primitive!) your jobs  
and get automatically the output retrieved when the  
jobs are finished

`crab.py -create all -submit all -monitor` to do all in just one  
command

If all is fine, present your work at meeting and offer some  
wine/beers/etc to crab team!

if problems, see next slides (and no beer for us...)



# Running CRAB

`crab.py -help` and read it!!!

`crab.py -create N` to create N jobs (no submission)

`crab.py -submit N -continue` to submit the job you have created  
(mind the `-continue`)

`crab.py -monitor -continue` to monitor (very primitive!) your jobs  
and get automatically the output retrieved when the  
jobs are finished

`crab.py -create all -submit all -monitor` to do all in just one  
command

If all is fine, present your work at meeting and offer some  
wine/beers/etc to crab team!

if problems, see next slides (and no beer for us...)



# Troubleshooting

- ▶ New born system: please be patient and provide feedback!

- ▶ most common problems:

`ConfigParser.ParsingError` error in `crab.cfg`: typically space at beginning of line!

Cannot create job type "ORCA" for VO "CMS" You did not eval 'scram runtime -csh'

Attempt to submit 1 jobs but only 0 jobs were created

`crab.py -submit 1` but not `-continue`

Need to create grid-proxy Create one `grid-proxy-init` dataset/owner not published with data tiers ... spelling

could be that dataset/owner available but not its parents (e.g. only DST and not Digi,Hit)



## Troubleshooting (II)

Job starts but ORCA crash many possibilities:

- ▶ your fault: **double check your code!!!**
- ▶ you are trying to access other data tiers (typically PU)
- ▶ problem with local Pool catalog:
  - ▶ rerun with: `CARFVerbosity=debug` and `PersistencyVerbosity=debug` and give the output to Site PubDB maintainer or savannah or `cms-wm-crab-feedback@cern.ch` list



# Troubleshooting (Job Aborted)

- ▶ You run `crab.py -mon -c` and you see **Aborted**. Many possibilities:
- ▶ Do **`edg-job-status <JodId>`**

Cannot plan: BrokerHelper: no compatible resources **today**  
ORCA deployed at site is 8\_7\_1, if newer, no  
matching resources can be found

Other messages send output of `edg-job-get-logging-info  
-v 2 <jobID>` to `support-eis@cern.ch`

