

Authentication and Authorization Management in Yabi

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Scientific Motivation for a Workflow Environment at IA2

GAPS (Global Architecture of Planetary Systems) is a long-term program for the comprehensive characterization of the architectural properties of planetary systems as a function of the hosts' characteristics (mass, metallicity, environment):

- more than 50 INAF and associated scientists in Italy, and from foreign institutes
- more than 10.000 spectra at TNG/HARPS-N since August 2012

GAPS required a strong interaction with HARPS-N private data

- Customizable re-process of GAPS data with appropriate spectral line mask and options;
- Perform queries on additional metadata content;
- A flexible and collaborative tool to manage additional info about the project and the observations;
- A repository where to access, synchronize, share and search for interesting data.



Yabi

- Yabi is a 3-tier application stack to provide users with an intuitive, easy to use, abstraction of compute and data environments. Developed at the Centre for Comparative Genomics and Murdoch University, Yabi has been deployed across a diverse set of scientific disciplines and high performance computing environments
- Yabi deployed at IA2 allows users to run custom data reduction with the HARPS-N DRS pipeline
- 10 tools, 8 different kind of workflows
- 28 custom masks
- Yabi access extended to all HARPS-N users with the same login credentials of the TNG archive (tools and masks developed by GAPS are only available to GAPS)
- More than 6000 workflows executed since March 2014

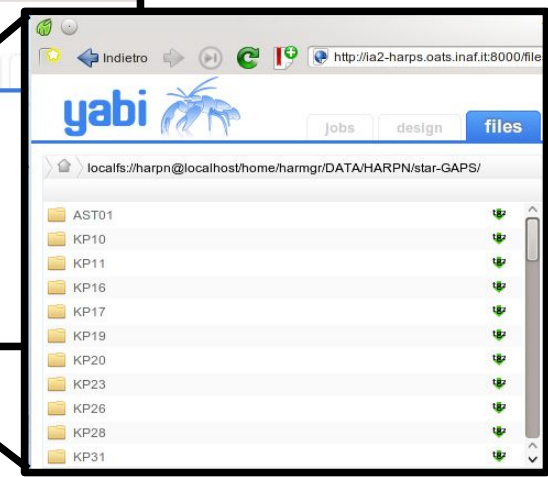


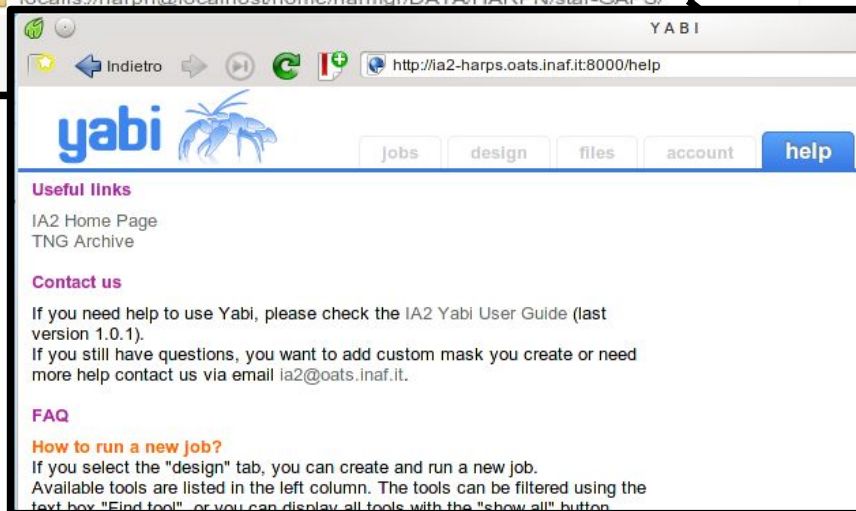
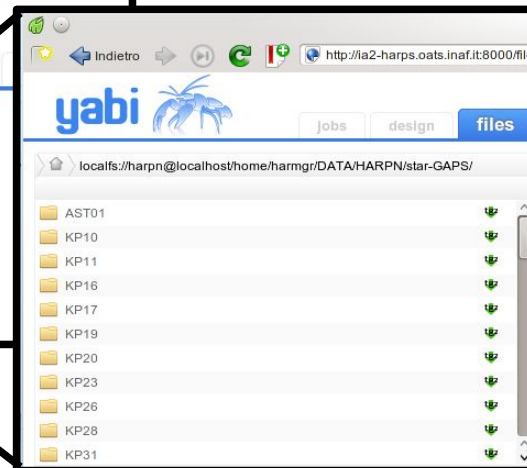
<https://ccgapps.com.au/yabi/>



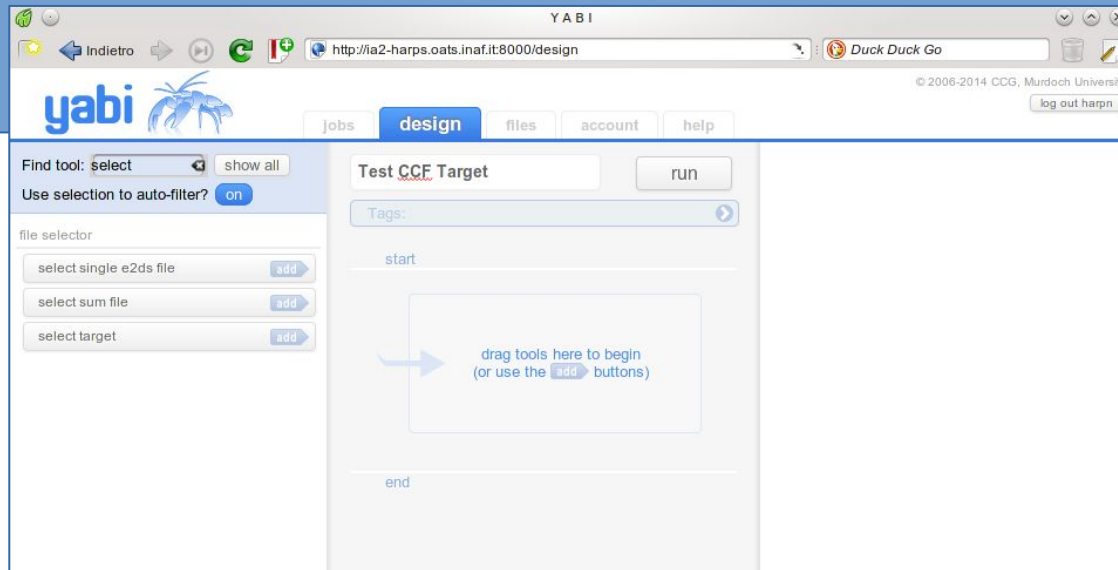
The screenshot shows a web browser window titled "YABI" with the address bar containing "http://ia2-harps.oats.inaf.it:8000/files". The page features a navigation menu with buttons for "jobs", "design", "files" (which is highlighted), "account", and "help". Below the menu is a file browser interface showing a list of directories:

- localfs://harpn@localhost/home/harmgr/DATA/HARPN/reduced-3.7.0-GAPS/
- localfs://harpn@localhost/home/harmgr/DATA/HARPN/reduced-3.6-GAPS/
- localfs://harpn@localhost/home/harmgr/DATA/HARPN/star-GAPS/



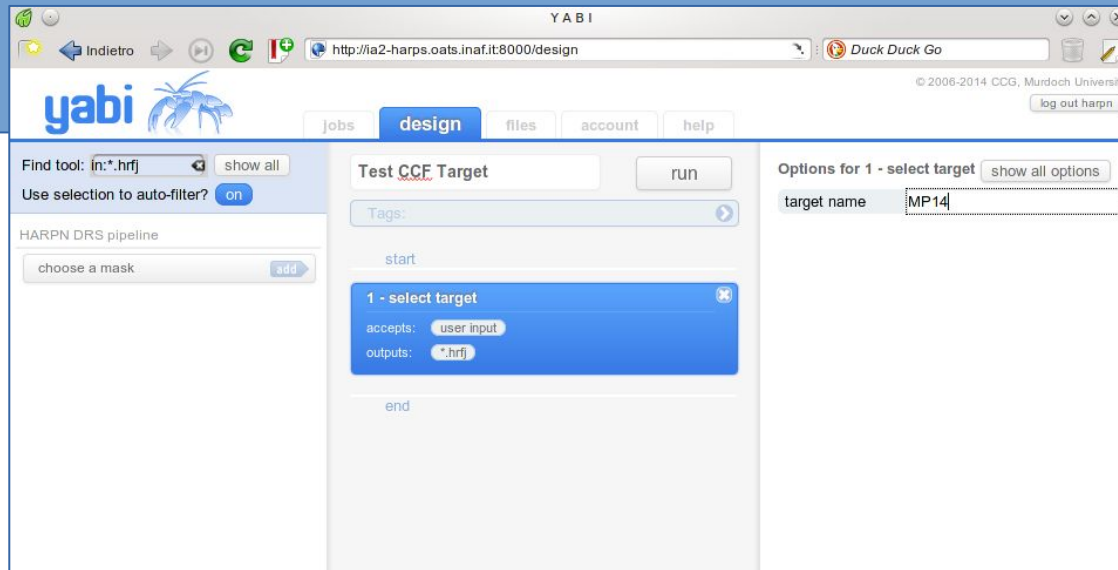


- From the “design tab” add a select tool to the workflow (e.g. select target)



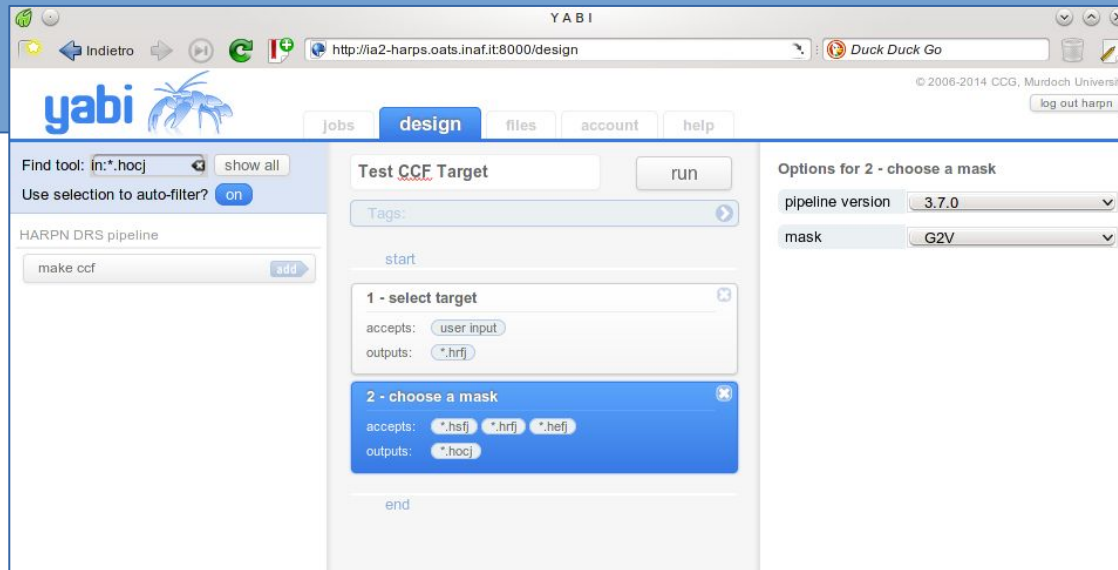
The screenshot shows the YABI web interface in a browser window. The address bar displays the URL `http://ia2-harps.oats.inaf.it:8000/design`. The page features a navigation menu with tabs for 'jobs', 'design' (which is active), 'files', 'account', and 'help'. A 'log out harpn' button is visible in the top right corner. The main content area is titled 'Test CCF Target' and includes a 'run' button. Below the title is a 'Tags:' field with a dropdown arrow. The workflow editor shows a 'start' node and an 'end' node, with a central box containing the instruction 'drag tools here to begin (or use the add buttons)'. On the left side, there is a 'file selector' panel with three options: 'select single e2ds file', 'select sum file', and 'select target', each with an 'add' button. Above the file selector, there is a 'Find tool: select' dropdown and a 'show all' button. Below the file selector, there is a 'Use selection to auto-filter?' checkbox which is currently checked and labeled 'on'.

- From the “design tab” add a select tool to the workflow (e.g. select target)
- Automatically Yabi suggests the next tool to add: choose a mask



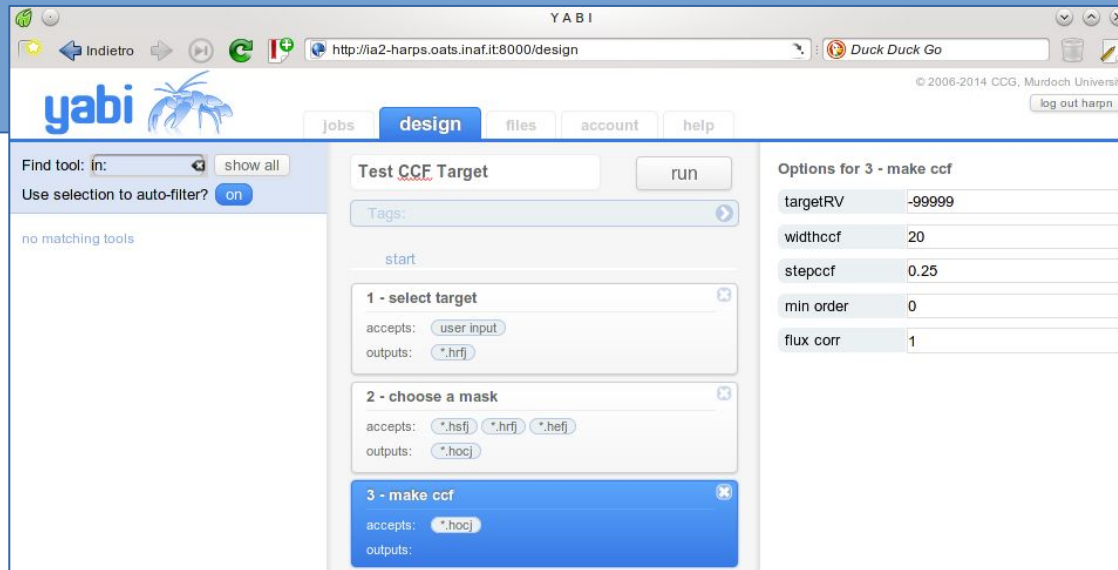
The screenshot shows the YABI web interface in a browser window. The address bar displays the URL `http://ia2-harps.ots.inaf.it:8000/design`. The page features a navigation menu with tabs for 'jobs', 'design', 'files', 'account', and 'help'. The 'design' tab is active, showing a workflow editor. On the left, a 'Find tool' search bar contains `in:*.hrfj` and a 'show all' button. Below it, a 'Use selection to auto-filter?' toggle is set to 'on'. A 'HARPN DRS pipeline' section includes a 'choose a mask' button with an 'add' icon. The main workflow area is titled 'Test CCF Target' and contains a 'run' button. A 'Tags' input field is present. The workflow steps are: 'start', '1 - select target' (highlighted in blue), and 'end'. The '1 - select target' step has 'accepts: user input' and 'outputs: *.hrfj'. To the right, an 'Options for 1 - select target' panel shows a 'target name' field with the value 'MP14' and a 'show all options' button. The footer of the browser window indicates '© 2006-2014 CCG, Murdoch University' and a 'log out harpn' button.

- From the “design tab” add a select tool to the workflow (e.g. select target)
- Automatically Yabi suggests the next tool to add: choose a mask
- Finally, add the tool “make ccf” and set the last input parameters



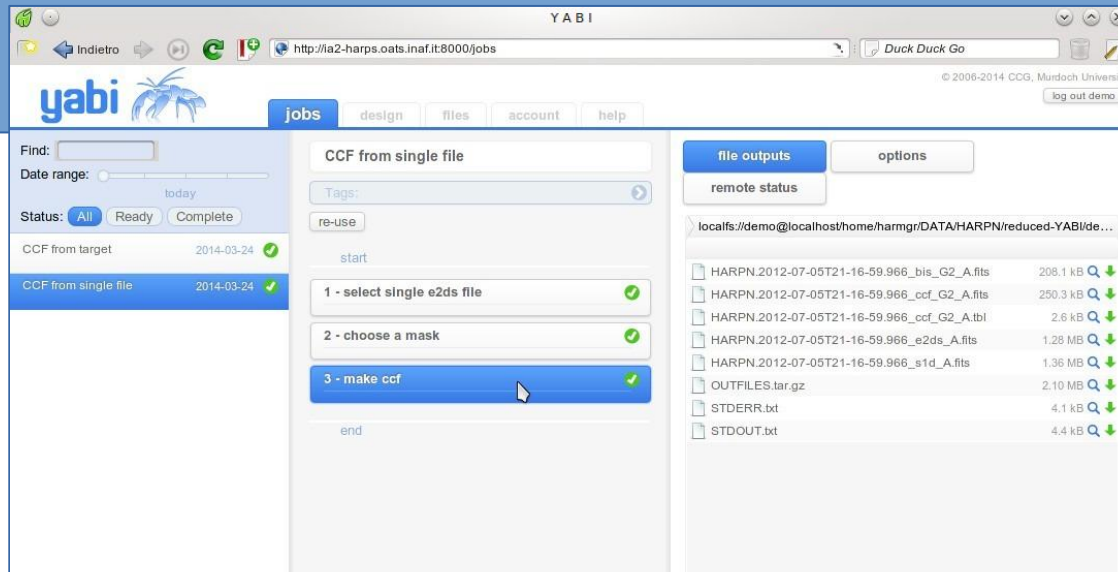
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- From the “design tab” add a select tool to the workflow (e.g. select target)
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- From the “design tab” add a select tool to the workflow (e.g. select target)
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The screenshot shows the YABI web interface. The browser address bar displays `http://ia2-harps.oats.inaf.it:8000/jobs`. The interface includes a navigation menu with tabs for 'jobs', 'design', 'files', 'account', and 'help'. A search bar is present at the top left. The main content area is titled 'CCF from single file' and shows a workflow with three steps: '1 - select single e2ds file', '2 - choose a mask', and '3 - make ccf'. The '3 - make ccf' step is highlighted in blue. To the right, there is a 'file outputs' section showing a list of files with their sizes and download links. The files listed are:

File Name	Size	Download
HARPN.2012-07-05T21-16-59.966_bis_G2_A.fits	208.1 kB	Download
HARPN.2012-07-05T21-16-59.966_ccf_G2_A.fits	250.3 kB	Download
HARPN.2012-07-05T21-16-59.966_ccf_G2_A.tbi	2.6 kB	Download
HARPN.2012-07-05T21-16-59.966_e2ds_A.fits	1.28 MB	Download
HARPN.2012-07-05T21-16-59.966_s1d_A.fits	1.36 MB	Download
OUTFILES.tar.gz	2.10 MB	Download
STDERR.txt	4.1 kB	Download
STDOUT.txt	4.4 kB	Download

Authorizations in Workflow Environment

- User access to data
 - User data
 - Archive data (remote)
 - Output data
- User access to computational resources
- User access to software (tools)
 - Open Software
 - Licenced software
 - Software developed by users

Backends, Credentials and Backend Credentials

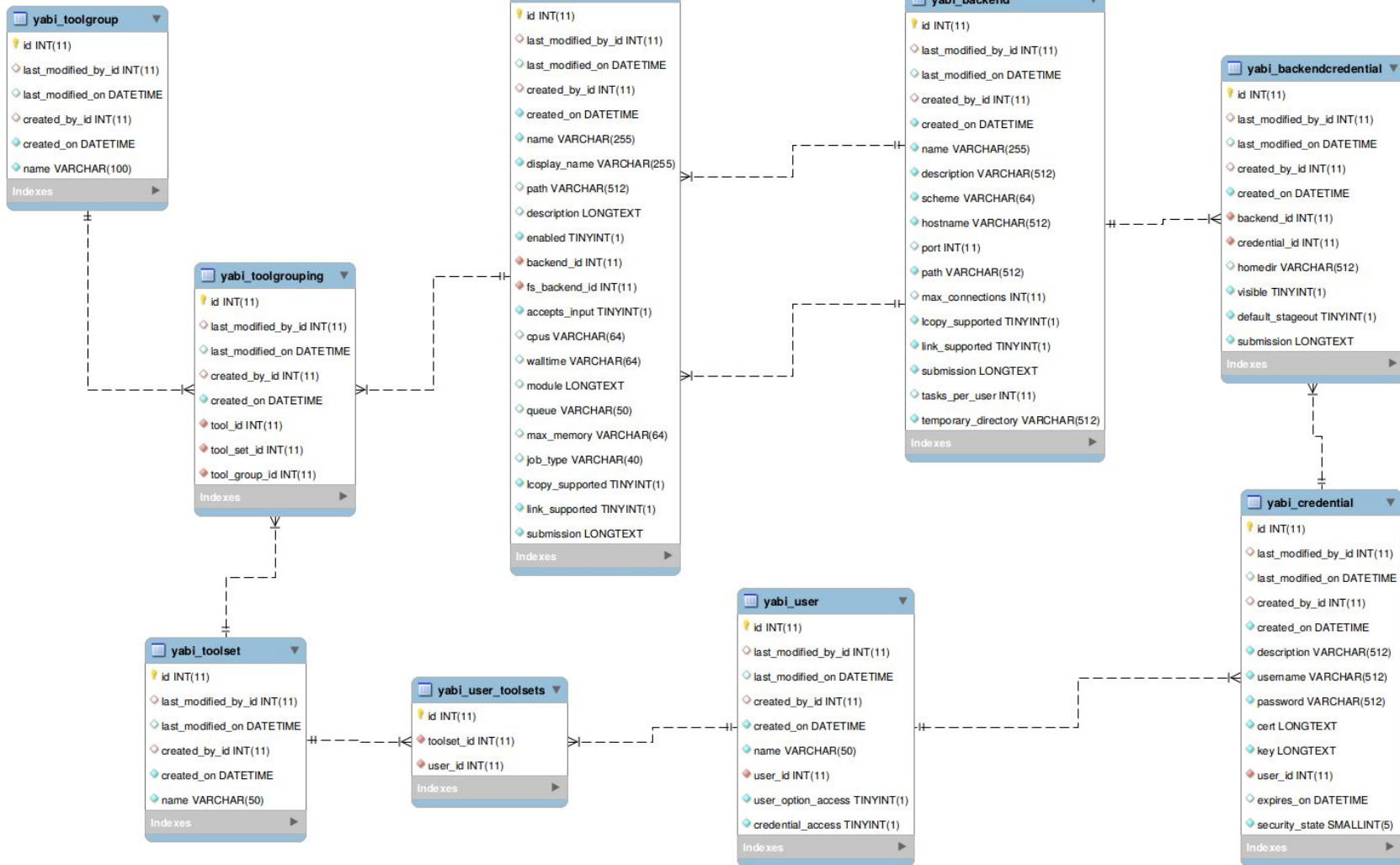
- The Yabi **backend** is a demon that provides execution and file services to the Yabi stack
- The Yabi backend abstracts away the details and complexity of individual protocols and resources to provide a consistent interface to the rest of Yabi
- Any tasks carried out by Yabi are always run as the user not as Yabi
- For each user you need to create a **credential** in order to allow that user to have access to backends (i.e. to execute jobs and workflows)
- To allow a user to use a backend, you need to connect the user credential to the backend through a **backend credential**

Backends, Credentials and Backend Credentials

- | | |
|--|---|
| <ul style="list-style-type: none">• In Yabi there are 3 types of Backend: execution, storage and null (for fileselector tools)• Several schema and connectors: localex, localfs, PBSPRO, SGE, Torque, SSH, SFTP, Slurm, Amazon S3, OpenStack Swift | <ul style="list-style-type: none">• A Credential is stored encrypted within the Yabi system and belongs to a Yabi user• In the Credential table all the fields will be needed, depending on the credential type: e.g. ssh key, certificate or user-passwd |
| <ul style="list-style-type: none">• Backend Credential is a linking table between a Credential and a Backend• Backend Credential also defines user's Default Stageout as a single directory where all the user's results will be staged out to | |

Tools, Toolsets, Toolgroups

- **Tools** are the elements the user builds the workflow with
- Yabi can run any command line tool installed on the machine where the tool will be executed
- Adding a new tool
 - A description of the tool and the parameters it accepts
 - Information about how to run the tool on a backend and the file system backend where outputs are saved
- Adding **toolsets** allows grouping of users to determine which tools they have access to
- **Toolgroups** determine how tools are grouped in the user interface
- Each tool in a **toolgroup** is assigned to a toolset, i.e. to a group of users that have access to that tool



Discussion Topics

- How to allow access to remote data resources?
- How to run workflows on data stored in geographically distributed sites?
- Standardize workflow description with Common Workflow Language (CWL)
- How to provide user access to perform custom data reduction on public data?
- What are the access policies for output data of workflows on public data?
- How to allow user to run their own software? How large this freedom should be?