\mathtt{q}_{py} Manual

\mathtt{q}_{py} The Queue Management System in Python

User, Administrator, and Developer manual

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1 For users

 q_{py} is used to submit jobs from a main server and execute them in different nodes, according the availability or the options given by the user. It can handle jobs running in both single and multi processors and it has several options to make a friendly user interaction. If you are using q_{py} for the first time, please read carefully the Section 1.1, Basics, just bellow. For a complete list of commands and their options, see the Section 1.2, Commands.

1.1 Basics

1.1.1 Installation

The main instalation will probably be done by your system administrator. You might have to do, however, the following thing:

Put the following lines in your ~/.bash_profile or ~/.bashrc:

```
1 export PATH=<qpy_dir>:$PATH
2 source <qpy_dir>/bash_completion.sh
```

Your system administrador should tell you which directory should be used in $\langle qpy_dir \rangle$. (S)He will might also send you a few files to be placed in your q_{py} directory, \sim /. qpy/. This directory has main configuration files, but in general you do not have to directly access them.

1.1.2 Initialisation

The first q_{py} command one should run is:

```
1 $ qpy restart
```

This command will set the background environment for using q_{py} . It should also be executed when some new update is done (you will be informed), or if the master node crashes.

1.1.3 Basic concepts and usage

Every job submmitted by q_{py} has an ID and a status. The ID, a fixed integer for a particular job, is unique and identifies your job in the pool of (your) jobs. The status can be:

- queue the job is waiting for allocation;
- running the was allocated and is running;

- done the job have run and is finished;
- killed the job had its execution killed by the user;
- undone the job have not been executed (it was killed while in the queue).

When you just submit a job, it has the queue status. When a node is allocated for it, it is executed and the status changes to running. If the job terminates normally, the status changes to done. If you kills the job (that is, stops its execution), the status changes to killed. If you kills a job that is still waiting for allocation, it will never be executed and the status changes to undone.

You can submit a job using the command sub:

```
1 $ qpy sub <job's execution command>
```

This means that you put, on your queue of jobs, a new job that has to be executed with the command $\langle \text{job's execution command} \rangle$. q_{py} will handle it for you from here on!

By the way, all commands in q_{py} are like this:

```
1 $ qpy <cmd> [options]
```

where $\langle cmd \rangle$ is one of the q_{py} commands and [options] are the otions for that command. Of course, these options depend on the actual command $\langle cmd \rangle$, some of them are optional, some not. See below for a complete list of commands and the explanation of their options.

Now that your job is on the list, you probably want to be able to:

- check the status of your job (or jobs);
- check the situation of the nodes;
- eventually kill an unsatisfactory job.

This can be done by the following commands:

```
1 $ qpy check
```

This shows a list of all your jobs. If you are interested only on a portion of these jobs, say, the ones that are currently running, try:

```
1 $ qpy check running
```

Another useful q_{py} command is:

```
1 $ qpy status
```

It will show all the available node, users, how many jobs each one is running, and a few extra information. If your jobs are waiting for too long on the queue, this command should provide an indication why.

Sometimes we are not so happy with the development of a job and we want to terminate its execution, without waiting to its normal termination. You can do this with:

```
1 $ qpy kill <job_ID>
```

where $\langle job_ID \rangle$ is the ID number of the job. Attention! This is not reversible. q_{py} will not ask you if you are sure to kill the job, so use the command with care.

There are several commands and options in q_{py} . Every command in q_{py} can be completed automatically by using the key <TAB>. It also gives the possible next arguments which, we hope, will be very helpful. Moreover, if you type:

```
1 $ qpy kill ? <TAB> <TAB>
```

where <TAB> <TAB> indicates that you should type the <TAB> key two times, you will receive a brief explanation on the command kill.

1.2 Commands

This section presents a list of all the commands and options of q_{py} . Remember that they should be used as:

```
1 $ qpy <cmd> [options]
```

where [options] depends on the command <cmd>.

1.2.1 restart

To start the q_{py} background environment. This command also works as an easy way to finish current session of q_{py} and start a new, which is most useful when the user wants to update the q_{py} to its latest version. If you are restarting q_{py} because the background environment has crashed, you might have to remove the file master_connection_port from the q_{py} directory before starting a new q_{py} session.

- Options: there are no options.
- Examples:

Restart the q_{py} background environment:

1 \$ qpy restart

1.2.2 finish

If you feel like done using q_{py} , please finish the existing q_{py} environment which is running in background using this command.

- Options: there are no options.
- Examples:

Finishes the background q_{py} environment:

```
1 $ qpy finish
```

1.2.3 sub

Used to submit any executable or command. The standard output and standard error of that command will be written in the files job_<job_ID>.out and job_<job_ID>.err respectively.

By default, q_{py} will allocate one core and 5 GB of RAM for such submission. To submit a job that uses more cores (say <N> cores) or a different amount of memory (say <M> GB), one can add this information in two ways. The first case is using the options -n <N> or -m <M> after qpy sub, but before the executable (or the options will be used as options to the command). The second case can be used when submitting a bash script as command. In this case, the following lines can be added in the script, that will be recognised by q_{py} :

```
1 #QPY n_cores=<N>
2 #QPY mem=<M>
```

- Options: the executable and its arguments; optionally, the the following flags *before* the executable:
 - -n <N> the number of cores
 - -m <M> requested memory, in GB
- Examples:

Executes the command hostname. Allocates the default one cores and 5 GB of memory for it:

```
1 $ qpy sub hostname
```

Executes the script ./script.sh (that must have the permission to be executed!). Allocates three cores and 10 GB of memory for it:

```
1 $ qpy sub -n 3 -m 10 ./script.sh
```

Executes the command 1s -ltr:

```
1 $ qpy sub ls -ltr
```

1.2.4 check

Checks the status of all the submitted jobs, printing a list of jobs. If no option is given, the command gives the list of all jobs submitted. One can also list just specific jobs, by their status, ID, or directory of submission. This information can be passed by the options, and if multiple options are given the jobs that satisfy any of the required property is listed. For example a status and a directory are given, this command will list all the jobs with that status plus the jobs that were submitted from the directory.

- Options: statuses, job IDs, and/or directories. The job IDs can be passed individually or as a range, such as 100-120.
- Examples:

Lists the jobs that are currently running:

```
1 $ qpy check running
```

Lists the jobs that have been submitted from the current directory

```
1 $ qpy check .
```

Lists the jobs with ID in the range 100-120, plus the jobs with ID in the range 130-135, plus the job 137:

```
1 $ qpy check 100-120 130-135,137
```

1.2.5 kill

Kills a particular job or set of jobs. One can kill several jobs by providing several job IDs (with a range, for instance, like in check. A status can be used to kill all jobs with that status (running or queue). Use all to kill all the jobs (be careful!).

- Options: one or more job IDs, a status (queue or running), or all.
- Examples:

Kills the jobs with ID 100, 101, 102, 103, 104, 105, and 108:

```
1 $ qpy kill 100-105 108
```

Kills all the jobs with status queue (that is, not yet allocated):

```
1 $ qpy kill queue
```

Kills all jobs:

```
1 $ qpy kill all
```

1.2.6 clean

Cleans the list of jobs that q_{py} has currently in memory. One can only clean jobs with status done, killed, or undone. Similarly to the kill command, the user can clean some specific jobs by the ID, all the jobs with some status, or all jobs. One can also clean by the directory of submission. It has no effect

- Options: one or more job IDs, a status (done, killed, or undone), all, or a directory.
- Examples:

Cleans the jobs with ID 100 to 200:

```
1 $ qpy clean 100-200
```

Cleans all the jobs with status undone (that ones that have been killed before running):

```
1 $ qpy clean undone
```

Cleans all the jobs that can be cleaned:

```
1 $ qpy clean all
```

1.2.7 status

This command is mainly useful for checking the multi-user environment. It writes the number of jobs running for each users, and in each of the available nodes.

- Options: There are no options.
- Examples:

```
1 $ qpy status
```

1.2.8 config

This command, if run without option, gives information about the current settings of q_{py} . Feel free to check, it is quite self explanatory. These settings can be changed by the same command, giving as option a keyword and a value (or soemtimes a sequence of values):

```
1 $ qpy config <keyword> <value> [<value2> ...]
```

In the following is a list of possible keywords and what they mean.

• colour true|false

The output of check is coloured (true) or not (false). Coloured output might not work for some terminals.

• coloursScheme <c queue> <c running> <c done> <c killed> <c undone>

Sets the colours to be used in a coloured output of check. Give always five colours, following the order above, that can be: grey, red, green, yellow, blue, magenta, cyan, and white.

checkFMT '<pattern>'

When the command check is used, the jobs are printed in a specific pattern. One can change this with this keyword, where pattern> is a string (the single quotes above are important!')
that may contain one of the "ampersanded" modifiers below, that will be replaced by the appropriate content of the job.

- $-\%j:job\ ID$
- -%s:job status
- -%c: command you used to submit the job

```
 %d : working directory of your job %n : node allocated for your job
```

- %N : number of cores for your job.

-%R: running time of the job. (time in queue if the job is in queue)

- %Q : actual time when the job is submitted

- %S : actual time when the job has started

- %E : actual time when the job has finished

It might seem a little complicated, but a example clarifies. The default value is:

```
%j (%s):%c (on %n; wd: %d)\n
```

Let us suppose that one has only one job, running a command bash my_script.sh, with ID 100, submitted from /home/user/, and it is currently running on the node comp1.

```
1 $ qpy check
2 100 (running):bash my_script.sh (on comp1; wd: /home/users/)
3 $ qpy config checkFMT '%j: %s\n'
4 Check pattern modified to '%j: %s\n'
5 $ qpy check
6 100: running
7 $ qpy config checkFMT '%j (%s)\n\tSubmitted from %d\n\tStarted at %S\n----\n'
8 Check pattern modified to '%j (%s)\n\tSubmitted from %d\n\tStarted at %S\n----\n'
9 $ qpy check
10 100 (running)
11
         Submitted from /home/users/
12
         Started at 2018-10-04 18:01:45.948201
13 ----
14 $ qpy config checkFMT default
15 Check pattern restored to the default value: '%j (%s):%c (on %n; wd: %d)\n'.
16 $ qpy check
17 100 (running):bash my_script.sh (on comp1; wd: /home/users/)
```

The \n is important, and it means "new line". Without it, all jobs are printed in the same line. One can also use \t to insert a tabulation, as the example shows.

• copyScripts <true|false>

Bash scripts, generally, stops if changes are made in it while it is still running. So it is sometimes interesting to have q_{py} running a copied script which allows to change and reuse the original bash script. If this keyword is set to true, qpy will 'qpy config' provides another advanced option to do this by copying the script to the local folder \sim /.qpy/scripts and use it later while running the actual calculation. This is very useful while working with a bash script. To use this, set it up with qpy config copyScripts true, where by default copyScripts is set to false.

The same thing can also be done for a specific job if the submitted script has the line #QPY cpScript true or if the job is submitted using the command qpy sub -c [script_name]. If the copyScripts is set True, then the reverse can be done by adding the line #QPY cpScript false in the script or by submitting the script using the command qpy sub -o [script_name].

- Options: a keyword and a value (or values) for this keyword.
- Examples: See above for examples for each keyword.

1.2.9 ctrlQueue

This command controls and move jobs in the queue. First the options pause and continue can be used to pause the submission of jobs from the queue to running, or to continue. Also, it can be used to move jobs within the queue (jump), and reorganise the order of submission.

• Options: pause, jump, continue

• Examples:

Consider the situation where you have a long list of jobs in the queue but you have some job(s) of top priority that needs to be done as soon as possible. The command ctrlQueue can be used in this situation to move the priority job(s) up in the queue. These are what you have to follow: The queue must first be paused, then "jump" the job <job ID> to <target>:

```
1     qpy ctrlQueue pause
2     qpy ctrlQueue jump <job ID> <target>
3     qpy ctlrQueue continue
```

The <target> can be a job ID, begin, or end.

1.2.10 tutorial

Shows a tutorial, that is basically a text version of the "For ssers" section of this manual.

- Options: Optionally, a keyword can be given to be searched in the tutorial
- Examples:

Opens the tutorial and goes to the first occurrence of ctrlQueue:

1 qpy tutorial ctrlQueue

2 For administrators

To work as administrator of q_{py} , it is useful to know a few facts about how q_{py} works. q_{py} has three levels:

- 1. the qpy
- 2. the qpy-master
- 3. the qpy-multiuser

The first, qpy, takes care of the interaction of the user with q_{py} , and its usage is described in the Section 1 of this manual. The second is the background environment of q_{py} of each user, and controls the user's jobs. The third is the global (or multi-user, as we prefer to say) background environment of q_{py} , that decides when and where a job can run, by allocating cores and memory to each q_{py} -master.

All this said, the duties of the administrator of q_{py} are:

- Maintain the program qpy-multiuser running;
- Add/remove new users and machines;
- Assist the users with any problems;
- Update q_{py} to newer versions.

This section explains how this is done. First of all, it is a good idea to have a user dedicated to the administration of q_{py} , what keeps everything more organised. We of course strongly discourage using the root for this task, for security reason. The administrator of q_{py} does not need any special permission.

2.1 Installation

We assume that you have a copy of q_{py} , otherwise you wouldn't be reading this manual. To set up a new q_{py} environment, the following steps must be taken:

- Make sure that every user that will use q_{py} has permission to read the directory where q_{py} is installed. This directory will be (and has been, in Section 1.1) called $\langle qpy_dir \rangle$.
- Create the configuration's directory in the administrator's home directory:

```
1 $ mkdir \sim/.qpy-multiuser/
```

• Create the following files in this directory:

```
1 $ vi ~/.qpy-multiuser/distribution_rules
2 $ vi ~/.qpy-multiuser/allowed_users
3 $ vi ~/.qpy-multiuser/nodes
4 $ vi ~/.qpy-multiuser/multiuser_connection_address
```

The content of these files are described in section 2.6. Please, read such section carefully and create these files according to your needs before proceeding.

• Finally, start the multiuser environment of q_{py} :

```
1 $ python <qpy_dir>/qpy-access-multiuser.py start
```

All the interaction of the administrator with q_{py} is made with the following command, plus some options:

```
1 $ python <qpy_dir>/qpy-access-multiuser.py
```

It is a good idea to define an alias for this by, for example, adding the following in the q_{py} administrator \sim /.bashrc:

```
1 alias qpy-admin='<qpy_dir>/qpy-access-multiuser.py'
```

2.2 Add new user

Let us suppose that a user with username $\langle user \rangle$ will start using q_{py} . As q_{py} usually runs across several machines, first make sure that $\langle user \rangle$ has the ssh keys properly configured, that is, that the user is be able to connect from the master node (where q_{py} is running) to the slave nodes without password. Because the home directory of the master node and the slave nodes must have the same home directory for a correct q_{py} environment, setting the ssh keys just once should be enough. To allow for the new user in q_{py} , add the username in the file $\sim /.qpy$ -multiuser/allowed_users, one user in each line:

```
1 $ cat ~/.qpy-multiuser/allowed_users
2 <user 1>
3 <user 2>
```

```
4 ...
5 <user>
```

After, you have to send the following files to the new user, to be put in his/her q_{py} directory. Something like this:

```
1 $ cp ~/.qpy-multiuser/multiuser_connection_address /home/<user>/.qpy/
2 $ cp ~/.qpy-multiuser/multiuser_connection_port /home/<user>/.qpy/
3 $ cp ~/.qpy-multiuser/multiuser_connection_conn_key /home/<user>/.qpy/
```

In addition, if you have a dedicated machine to run q_{py} , copy also the following file file to the user's q_{py} directory:

```
1 $ cat ~/.qpy-multiuser/master_connection_address
2 <hostname>
3 $ cp ~/.qpy-multiuser/master_connection_address /home/<user>/.qpy/
```

If this file is not present in the user's q_{py} directory, q_{py} runs locally.

2.3 Add and remove a node

To add or remove a node, edit the file \sim /.qpy-multiuser/nodes (see Section 2.6), and run the following commands:

```
1 $ python <qpy_dir>/qpy-access-multiuser.py nodes
2 $ python <qpy_dir>/qpy-access-multiuser.py distribute
```

2.4 Update version

When a new version of q_{py} is released, depending where the new changes have been done, you might have to restart the multiuser background environment:

```
1 $ python <qpy_dir>/qpy-access-multiuser.py finish
2 $ python <qpy_dir>/qpy-access-multiuser.py start
```

And/or ask all the users to restart their own masters, with:

```
1 $ qpy restart
```

Whatever is the case, this will be informed in the release's notes.

2.5 Commands

This is a full list of the commands that are available to the administrator:

• start

Starts a new qpy-multiuser instance. Run this if you are starting q_{py} for the first time, if a update is needed or if the machine qhere q_{py} runs has crashed.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py start
```

• finish

Finishes the multiuser background environment.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py finish
```

• nodes

Loads the content of the file \sim /.qpy-multiuser/nodes.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py nodes
```

• distribute

Distributes the cores among the users.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py distribute
```

• variables

Lists several internal variables of qpy-multiuser. Used mainly for debugging.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py variables
```

• status

Show the current status of the users, nodes and cores. It is the same as the command status accessible to the users.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py status
```

• saveMessages

Saves messages from the internals of qpy-multiuser, that will be shown in the "variables" command. Mainly for debugging.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py saveMessages
```

There are some "cheatting" commands that the administrator can run, such as artificially adding or removing running jobs of users. These commands are not supposed to be used in a normal run, but only if something went wrong and must be manually fixed. Use only if you know exactly what you are doing!!

• __user

Adds a new user to apy-multiuser. In a normal run, it is automatically done when the user restart his/her q_{py} .

```
1 $ python <qpy_dir>/qpy-access-multiuser.py __user <user_name> \
2 <address> <port> <conn_key>
```

• __req_core

Asks for a slot (cores plus memory) to run a job. In a normal run, it is done by the user's master whenever there is a job in his/her queue.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py __req_core <user_name> \
2 <jobID> <n_cores> <mem> <queue_size>
```

__remove_job

Tells the multiuser background environment that a job has finished. In a normal run, it is done by the user's master whenever the job has finished.

```
1 $ python <qpy_dir>/qpy-access-multiuser.py __remove_job <user_name> \ 2 <job_ID> <queue_size>
```

2.6 Files

This is a list of the files in the qpy-multiuser directory. Some of them the administrator should edit to control the behaviour of q_{py} , some others not.

• distribution_rules

This file defines how the cores are distributed to the users. The basic syntax is one of the following:

```
1 even minimum <n_cores>
```

This means an even distribution among the users, with at least <n_cores> granted for each.

allowed_users

A list with all the users that can use the q_{py} environment:

```
1 <user_1>
2 <user_2>
3 <user_3>
```

• nodes

A list with all the nodes available in the q_{py} environment. Each line has the information of one node, as shown below:

```
1 <node_1> <n_cores> [M]
2 <node_2> <n_cores> [M]
3 <node_3> <n_cores> [M]
```

First is the hostname of the node, followed by the number of cores this node has (or is available to q_{py}) and, optionally, a "M", to indicate that that node has preference for multicore jobs.

multiuser_connection_address

This file simply contains the address where the qpy-multiuser instance will run. If you do not set this, q_{py} automatically uses the local machine (localhost).

```
1 <hostname for qpy-multiuser>
```

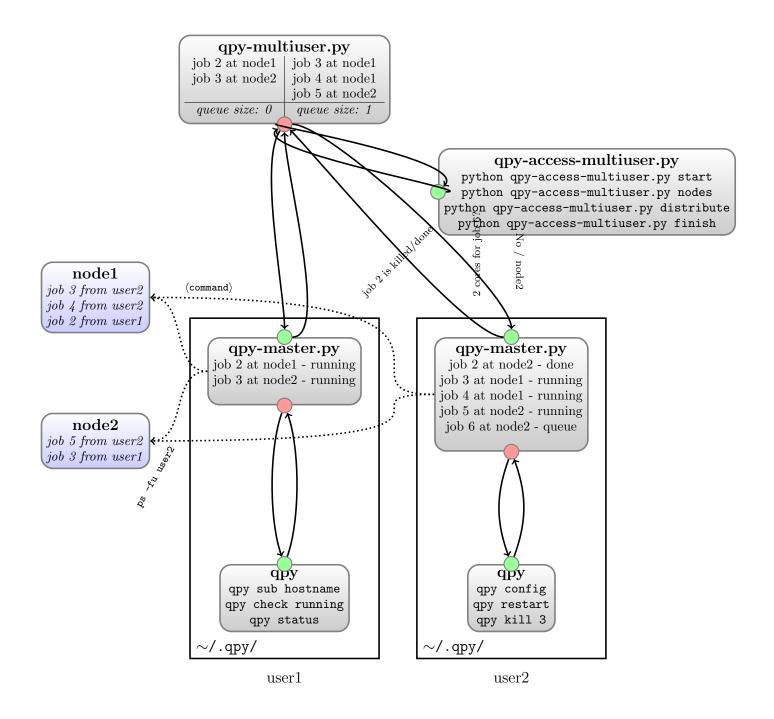
Optionally, for the sake of organisation mainly, the programs qpy-multiuser and the qpy-master of each user can run on a machine different than the node where the users work and submit their jobs from. For instance, you might have a machine (even a virtual machine) dedicated to qpy. To do this, <nostname for qpy-multiuser> must be the hostname of this machine In this case, the files master_connection_address that the users have in their qpy directory should have the correct hostname.

The following files are generated and used by q_{py} , but the administrator should not alter or delete them. Moreover, these files should not be shared, because they have the information required to make

all the message transfers. q_{py} takes care of the permission of the directories \sim /.qpy-multiuser/ and \sim /.qpy/ of each user (whose content should also not be shared).

- $\bullet \ \, \sim\!\!/\,.\, \texttt{qpy-multiuser/multiuser_connection_port}$
- $\bullet \ \, \sim\!\!/\,.\, \texttt{qpy-multiuser/multiuser_connection_key}$
- \bullet $\sim\!\!/.qpy\mbox{-multiuser/connection_<user>_key}$
- \bullet $\sim\!$ /.qpy-multiuser/connection_<user>_key
- $\bullet \ \, \sim\!\!/\,.\, \texttt{qpy-multiuser/connection_<user>_key}$

3 For developers



Python's multiprocessing: Listener/Client connection

