How to run a simple serial job?

A simple submission is

```
[testuser@login-node-1-0 basic-test]$ qsub simple_serial_job.txt
### where the script looks like this
[testuser@login-node-1-0 basic-test]$ cat simple_serial_job.txt
#$ -q all.q    ### specify which queue you want
#$ -cwd        ### not necessary but useful; it tells the system you want to start
### from the same folder you were when you submitted the job
./myprogram.x
```

Please check out the explanation of the policy and the list of queues to decide where to submit your calculations.

To submit in a lab-specific queue you will need to modify the script in a way similar to this (refer to the list of available queues for specific information)

```
[testuser@login-node-1-0 basic-test]$ cat simple_serial_job_lab_queue.txt
#$ -q mylab.q    ### specify which queue you want
#$ -cwd          ### not necessary but useful; it tells the system you want to start
### from the same folder you were when you submitted the job
./myprogram.x
```