

# **P56 - Lab 2, Measuring: Disks and Networks**

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Lent 2023/2024

The goal of this lab is to gain experience measuring two computer sub-systems: two of the five principle contributors to (poor) systems performance. You will tackle a seemingly straightforward problem, estimating the time each will approach, refine your approach, and interpret the measurements you have made.

As part of this lab you will be following the P56 textbook, you are keenly encouraged to record your lab progress; you may wish to use a blank lab notepad, or a Jupiter notebook to record your results, annotating with figures as appropriate.

This lab will require you build and extend a measurement-focussed codebase. The code base can be retrieved from <https://www.cl.cam.ac.uk/teaching/current/P56/ucamonly/book-user-code.zip>

We will closely follow Chapter 5 and 6 detailing the measurement of a disk system and of one side of a network client-server.

## **1 Before you begin**

Update your pull of the P56 repository.

Textbook tools create html viewable using a browser; to extract svg files (e.g., for your lab reports) you can use the tool SVG crowbar 2 from <https://nytimes.github.io/svg-crowbar/>

## **2 Notes for Chapter 5 - Disk**

Follow the chapter closely; including the Exercises 5.1 to 5.11 as appropriate.

On each R.Pi there are two disks the root is an sd-card, while /flash/ is a USB thumb drive based upon classic flash memory.

Furthermore, on the first machine of each cluster is a faster drive, an SSD disk. It is mounted at /nvme

There are a limited number of HDD disks with ye-olde spinning platters. Two machines in the cluster:

l51-pi046.nf.cl.cam.ac.uk 5400RPM 160GB mounted as /hdd5400

and

l51-pi047.nf.cl.cam.ac.uk 7200RPM 500GB mounted as /hdd7200

Your l51 login credentials will permit **all of you** access to each of these two machines but beware you are sharing these machines with your colleagues so coordinate usage, use unique directory names (crsid) and play nicely please.

Compile the mystery3 code with the command-line:

```
g++ -O2 -o mystery3 mystery3.cc -lrt
```

Note that this code tries to strip the extension name naïvely, looking for the last dot. This means that the path to use must either contains a file that has an extension name, or it should not include any dot. E.g. ../../nvme\_testfile will cause problems, but 2 /nvme/speedTestFile will not.

When Processing, for the sort make sure

```
export LC_ALL=C
```

is set (otherwise the sort will not function predictably.)

An alternative approach is in this example

```
cat testfile_read_times.json | LC_ALL=C sort | \
./makeself show_disk.html > testfile_disk_read.html
```

This puts LC\_ALL=C near sort to avoid changing environment variables globally.

Programs such `makeself` expect to be run in the `book-user-code` directory.

The html files may be viewed using the browser running on the R.Pi

```
chromium-browser testfile_disk_read.html testfile_disk_write.html
```

You can also copy the html files locally for faster response.

### Note Well

As indicated in Exercise 5.9 you will need to write some code in `mystery3.cc`;

5.9 Complete the missing part of `TimeDiskWrite()`. Mine is seven more lines, setting the block current times. This will be easy if you have followed what the strategy is, and a bit harder if you have been only skimming this text and the code. But when you are done, you will better understand what is going on.

Search in `mystery3.cc` for "You get to fill in this part !!"

Don't forget to recompile and if you get it wrong (or the code doesn't exist) the write graph will appear to have taken no time at all – a vertical line on the graph in `testfile_disk_write.html`

## 3 Notes for Chapter 6 – Networks

Like the previous, follow Chapter 6 and attempt the exercises at the end: 6-1 to 6-3

While no code needs to be written; like the code of Chapter 5, the executables that process data expect to be run in the `book-user-code` directory. You will need two machines for this chapter so will need to have the code (or at least the server executable) on each machine too.

A few typo's remain in the textbook, former TA Gary Guo has provided some example commands on the web page <https://hackmd.io/@nbdd0121/ByAvCuMai>. I've compiled corrections from his examples and others below:

Compiled with `./compile_all_user.sh` (alternatively Gary supplied a makefile, posted on his website above).

On your\_server\_machine

```
./server4
```

On your\_client\_machine

```
./client4 your_server_machine 12345 write -key "kkkkk" -value "vvvvv" 1000000
```

```
./client4 your_server_machine 12345 quit
```

Each time you run `client4`, it will display the log filename it writes to. You need only remove `.log` and assign it to a var for convenience. Each command will write a new logfile; you are interested in the log from the write command, not the quit command.

For example

```
LOGSTEM=client4\_20230209\_141229\_151-pi888.nf.cl.cam.ac.uk\_19102
```

Processing:

```
./dumpplogfile4 "Write 1MB" $LOGSTEM.log > $LOGSTEM.json  
cat $LOGSTEM.json | ./makeself show_rpc.html > $LOGSTEM.html
```

Once again you can display the html files using chromium on the R.Pi, or copy them locally.

## 4 Saving Your Experiments

Make sure to back up your experiments, including (but not limited to) Jupyter notebooks, dump files and scripts. Remember that multiple teams may use the same test machines, so be careful when handling data.

All the measurements are saved under your crsid folder, so backing up the entire folder is a good idea. To copy a remote directory onto your local machine:

`sftp 151@<hostname>.nf.cl.cam.ac.uk` and `get -r <directory>`.

There are also other ways to copy a remote directory, you are welcome to use those as well. You may wish to compress results files in order to save space.

Exporting a Notebook as `.tex` will save graphs as separate files, which you can then include in your lab report.

Please do not push any changes, data or results directly to the P56 repository. You can fork the repository to your own user and push changes there. If you would like to suggest a correction or an enhancement to a notebook or a script, please use pull-requests.

## 5 Understanding Your Measurements

A single lab report will be required for the first three labs. Instructions for the lab report will be provided separately.

The following items are intended to help you understand your results, and may provide supporting evidence for your report. However, they are just suggestions - feel free to approach the data differently!

- Discuss the methodology of the measurement tools.
- Explain how the limitations of these are mitigated in this lab.
- Explore the limitations of the experiments conducted in this lab, and explain where the quality of the experiment (e.g., setup, methodology) could have been improved.

You should always look for odd or surprising results, and try to explain them. Note that sometimes exceptional results may indicate a problem in your setup or scripts.